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Sequence 44, Appl
Sequence 42, Appl
Sequence 91, Appl
Sequence 51, Appl
                                       January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                     Sequence 43,
                                                                                                                                                                                                                                                                                                                                                                        Description
                                                                                                                                                                                GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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1 US-10-354-240-44

1 US-10-354-240-42

1 US-10-354-250-42

1 US-09-994-595-91

1 US-09-813-33-51

1 US-09-813-135-103-51

1 US-08-424-5508-366

1 US-10-256-850-35

1 US-09-826-290-39

1 US-09-826-290-31

1 US-09-826-290-415
                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                              1608061 segs, 361289386 residues
                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                             protein search, using sw model
                                                                              1 GVVPVHAQDGDAITMRNVTD 20
                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                  US-09-202-464-17
104
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Match Length
                                                                                                                                       length: 0
length: 20
                                                                                                                                                                                                                                                                                                                                                                                          Minimum DB seq
Maximum DB seq
                                                                  Title:
Perfect score:
                                                                                          Scoring table:
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                                                                               Sequence:
                                                                                                              Searched:
                                                                                                                                                                                   Database
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No.
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Sequence 43, Application US/10354240

Sequence 43, Application WS/10354240

Publication No. US2003018847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Kine, Akinori

APPLICANT: Mama, Akiko

FILE REFERENCE: SPO-103D1

CURRENT PELINGATION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Diserring APPLICANT:

FILE REFERENCE: SPO-103D1

CURRENT PELINGATION NUMBER: US/10/354,240

CURRENT FILING DATE: 1997-03-10

FRIOR FILING DATE: 1997-03-10

PRIOR FILING DATE: 1998-09-09

WUMBER OF SEQ ID NOS: 174

SEQ ID NO 43

SEQ ID NO 43
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                                             Sequence 57, Appl
Sequence 45, Appl
Sequence 248, Appl
Sequence 378, Appl
Sequence 378, Appl
Sequence 248, Appl
Sequence 248, Appl
Sequence 248, Appl
Sequence 378, Appl
Sequence 378, Appl
Sequence 378, Appl
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Sequence 12, A
Sequence 21, A
Sequence 672, A
Sequence 675, A
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Sequence 3
Sequence 3
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Sequence
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Pred. No. 0.0023;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (1). (15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 29
US-10-264-309-190
US-10-256-850-56
US-10-256-850-56
US-10-256-850-57
US-09-735-705-248
US-09-735-705-248
US-09-850-716A-248
US-09-850-716A-248
US-09-850-716A-248
US-09-897-778-248
US-09-897-778-248
US-09-897-778-248
US-10-117-982-248
US-10-117-982-248
US-10-117-982-378
US-10-117-982-378
US-10-313-986-248
US-10-356-191-12
US-09-999-186-675
US-09-999-186-675
US-09-999-186-675
US-09-999-186-675
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Best Local Similarity 73.3%; Pr
Matches 11; Conservative 2;
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ORGANISM: Cryptomeria japonica
      NAME/KEY: MISC FEATURE LOCATION: (1)..(15)
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1 GVVPVHAQDGDAIIM 15

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Sequence 39, Appl Sequence 61, Appl Sequence 415, App

Sequence 53,

Sequence 51, 1 Sequence 51, 1 Sequence 366, 1 Sequence 13, 1

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; Sequence 51, Application US/0981333
; Patent No. US2002011916041
; GENERAL INFORMATION:
    APPLICANT: DeGroot, Anne S
    TITLE OF INVENTION: Human T Cell Response to MHC-Binding Motif Clusters
; TITLE REFERENCE: 17999-004 US
; CURRENT APPLICATION NUMBER: US/09/813,333
CURRENT APPLICATION NUMBER: 60/190,834
PRIOR PLING DATE: 2000-03-20
; NUMBER OF SEQ ID NOS: 81
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 51
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bhattacharjee, J.
APPLICANT: Bhattacharjee, J.
APPLICANT: Brattacharjee, J.
APPLICANT: Brattacharjee, Vasker
TITLE OF INVENTION: METHODS AND REAGENTS FOR DETECTING FUNGAL PATHOGENS IN
TITLE OF INVENTION: A BIOLOGICAL SAMPLE
FILE REPERENCE: 96.247-A
TITLE OF INVENTION: A BIOLOGICAL SAMPLE
FILE REPERENCE: 96.247-A
CURRENT APPLICATION NUMBER: 08/650,809
PRIOR FILING DATE: 1997-05-20
NUMBER OF SEQ ID NOS: 160
SOFTHARE: Microsoft Word 97
SEQ ID NO 91
LENGTH: 19
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                            Gaps
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Pred. No. 44;
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Pred. No. 3.3;
0; Mismatches
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; ORGANISM: Mycobacterium tuberculosis
US-09-813-333-51
                                                                                                                                                                                                                                                                                                       Sequence 91, Application US/09994595
Publication No. US20030039981A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 35.6%;
Best Local Similarity 66.7%;
Matches 6; Conservative
Similarity 80.0%;
8; Conservative
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                                                                                            1 GVVPVHAQDG 10
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Best Local Similarity
Matches 8; Conserva
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Best Local Similarity
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12 VVPLHRSDG
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US-10-044-703-51
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PRIOR FILING DATE: 1998-09-09 NUMBER PRIOR FILING PRIOR FILING DATE: 1998-09-09 NUMBER PRIOR FILING DATE: 1998-09-09
                                                                                                                                                                                                                                                                                 APPLICANT: Kume, Athori
APPLICANT: Kume, Athori
APPLICANT: Kume, Athori
APPLICANT: Name, Akico
APPLICANT: Dairiki, Kazuo
APPLICANT: Imame, Akico
APPLICANT: Imame, Akico
APPLICANT: Imame, Akico
APPLICANT: Imame, Akico
APPLICANT: STO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: DS/142,524
PRIOR FILING DATE: 1998-99-99
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 44
LENGTH: 15
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, OTHER INFORMATION: Cryjl peptide, Figure 1, Row 30 US-10-354-240-44
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INFORMATION: Cryjl peptide, Figure 1, Row
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Pred. No. 0.049;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 42, Application US/10354240 Publication No. US20030185847A1
                                                                                                                                                                     Sequence 44, Application US/10354240 Publication No. US20030185847A1
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ORGANISM: Cryptomeria japonica
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ORGANISM: Cryptomeria japonica
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Best Local Similarity 60.0%;
Matches 9; Conservative
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                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Sone, Toshio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MISC_FEATURE
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LENGTH: 15
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FILING DATE:
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Fublication No. US20040057961A1

GENERAL INFORMATION:

APPLICANT: Brown University Research Foundation

APPLICANT: Brown University Research Foundation

APPLICANT: Brown University Research Foundation

APPLICANT: DeGroot, Anne S

TITLE OF INVENTION: Human T Cell Response to MHC-Binding Motif Clusters

FILE REPERENCE: 17999-004-061

CURRENT APPLICATION NUMBER: 09/813,333

PRIOR APPLICATION NUMBER: 09/813,333

PRIOR FILING DATE: 2000-03-20

PRIOR FILING DATE: 2000-03-20

NUMBER OF SEQ ID NOS: 81

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 51

LENGTH: 20
Sequence 51, Application US/10044703
GENDICALION NO. US20020192233A1
GENDICAL INFORMATION.
APPLICANT: DeGroot, Anne S
TILLE OF INVENTION: Human T Cell Response to MHC-Binding Motif Clusters
FILE REFERENCE: 17999-004 US
CURRENT APPLICATION NUMBER: US/10/044,703
CURRENT FILING DATE: 2002-05-20
PRIOR PAPLICATION NUMBER: 60/190,834
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 81
SSOFTWARE: Patentin Ver. 2.1
LENGTH: 20
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TAMI J. PILOT-MATIAS
GEORGE J. DAWSON
GEORGE G. SCHLAUDER
SURESH M. DESAI
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BUIJK
                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Mycobacterium tuberculosis
US-10-044-703-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT; ORGANISM: Mycobacterium tuberculosis
US-10-239-103-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 366, Application US/08424550B Publication No. US20020119447A1
                                                                                                                                                                                                                                                                                                                                                                    Query Match 33.7
Best Local Similarity 66.7
Matches 6; Conservative
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Matches 6, Conservative
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12 VVPLHRSDG 20
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APPLICANT: JOHN N
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US-08-424-550B-366
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APPLICANT:
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APPLICANT:
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APPLICANT:
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APPLICANT:
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APPLICANT: Irimura, Tatsuro
APPLICANT: Irimura, Tatsuro
APPLICANT: Irimura, Tatsuro
APPLICANT: Wateumoco, Mariko
APPLICANT: Win, Mijung
APPLICANT: Yin, Mijung
APPLICANT: Or Takashi
TITLE OF INVENTION: Lectins for Analyzing Sugar Chains and Method of Using the Same
FILE REPERENCE: 03-786
CURRENT APPLICATION NUMBER: US/10/468,543
CURRENT FILING DATE: 2001-08-20
BRIOR FILING DATE: 2001-02-20
NUMBER OF SEQ ID NOS: 95
SOFTWARE: Patentin Version 3.1
LENGTH: 11
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; OTHER INFORMATION: Generated from randomly recombinant DNA part of MAH.
US-10-468-543-13
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APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 116
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 32; DB 8; L
Pred. No. 2.4e+02;
                                                                                                              ADDRESSEE: ABBOTT LABORATORIES D377/AP6D STREET: 100 ABBOTT PARK ROAD CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                      COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435435
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6355
TELEPAX: 708-938-2623
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100.0%; Pre
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 366: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16 amino acida
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MOLECULE TYPE: protein
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ORGANISM: Artificial
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Best Local Similarity
Matches 6; Conserv
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4 VPVHAQ 9
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APPLICANT: White, Front
APPLICANT: White, Front
APPLICANT: White, Front
APPLICANT: White, Front
TITLE OF INVENTION: Nucleic Acid Molecules, Polypeptides and
TITLE OF INVENTION: Uses Therefor, Including Diagnosis and Treatment of
TITLE OF INVENTION: Alzheimer's Disease
FILE REFERENCE: 2572-1-001 N2
CURRENT APPLICATION NUMBER: US/09/0826,290
CURRENT APPLICATION NUMBER: US 60/194,504
PRIOR APPLICATION NUMBER: US 60/194,504
PRIOR PILING DATE: 2000-04-30
PRIOR PILING DATE: 2000-011-28
NUMBER OF SEQ ID NOS: 492
SOFTWARE: FastSEQ for Windows Version 4.0
SEG ID NO 61
LENGTH: 14
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Sequence 415. Application US/09826290
Patent No. US20020146668A1
GENERAL INFORMATION:
APPLICANT: Dutham, L.Kathryn
APPLICANT: Friedman, David L.
APPLICANT: Friedman, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Ammel, Lida H.
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Durham, L.Kathryn
APPLICANT: Durham, L.Kathryn
APPLICANT: Friedman, David L.
APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Kimmel, Lida H.
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Potter, David M.
APPLICANT: Solff, Christian
APPLICANT: Silber, B. Michael
APPLICANT: Silber, B. Michael
APPLICANT: Sunderland, P. Trey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31; DB 9; Length 14;
Pred. No. 3.1e+02;
1; Mismatches 5; Indels
                                                                                                                                                             Length 14;
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Pred. No. 3.1e+02;
1; Mismatches 5; Indels
NUMBER OF SEQ ID NOS: 492
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 39
LENGTH: 14
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Patent No. US20020164668A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Townsend, Robert Reid
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Best Local Similarity 50.0
Matches 6; Conservative
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; ORGANISM: homo sapien
US-09-826-290-61
                                                                               TYPE: PRT

CRGANISM: homo sapien

US-09-826-290-39
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Best Local Similarity
Matches 6; Conserv
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APPLICANT:
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                                                                                                                                 Sequence 53, Application US/10256850
; Sequence 53, Application US/10256850
; Publication No. US20040024178A1
; GENERAL INFORMATION:
APPLICANT: ASHMAN, STEPHEN
APPLICANT: BLACK, MICHAEL T.
APPLICANT: BROTTON, GORDON
APPLICANT: MOONE, KEITH JAMES
TITLE OF INVENTION: BACTERIAL SIGNAL PEPTIDASE INHIBITORS;
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: P32237
CURRENT FILING DATE: 2002-09-27
PRIOR APPLICATION NUMBER: US/10/256,850
CURRENT FILING DATE: 2001-08-03
PRIOR PAPLICATION NUMBER: PCT/EP00/00751
PRIOR PAPLICATION NUMBER: GB 9902399.6
PRIOR PILING DATE: 2001-08-03
PRIOR PILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 64
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 53
LENGTH: 13
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Pred. No. 2.8e+02;
1; Mismatches 2; Indels
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US-10-256-850-53
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Best Local Similarity 62.5%;
Matches 5; Conservative
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3 LPAHAADG 10
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        2 VVPVHAQD 9
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2 VMPLHLQD 9
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APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES, POLYPEPTIDES AND USES THEREFOR,
TITLE OF INVENTION: INCLUDING DIAGNOSIS AND TREATMENT OF ALZHEIMER'S DISEASE
FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
PRIOR FILING DATE: 2001-10-03
                                                APPLICANT: Sunderland, P. Trey
APPLICANT: Sunderland, Robert Reid
APPLICANT: Townsend, Robert Reid
APPLICANT: Townsend, Robert Reid
APPLICANT: Williams, Stephen A.
TITLE OF INVENTION: Uses Therefor, Including Diagnosis and Treatment of
TITLE OF INVENTION: Alzheimer's Disease
FILE REFERENCE: 2572-1-001 N2
CURRENT APPLICATION NUMBER: US/09/826,290
CURRENT APPLICATION NUMBER: US 60/194,504
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/253,647
PRIOR FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 492
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 415
LENGTH: LAND
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Pred. No. 3.1e+02;
1; Mismatches 5; Indels
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Publication No. US20040022794A1
GENERAL INFORMATION:
                 Silber, B. Michael
Stiger, Thomas R.
Sunderland, P. Trey
Townsend, Robert Reid
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SNYDER, PETER J.
SOARES, HOLLY D.
STIGER, THOMAS R.
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SOFTWARE: Patentin Version 2.1
SEQ ID NO 190
LENGTH: 14
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Rohlff, Christian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: DURHAM, L. KATHRYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAJESH B.
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Best Local Similarity 50.0
Matches 6; Conservative
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; ORGANISM: Homo sapiens
US-10-264-309-190
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POTTER,
ROHLFF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SILBER,
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being printed,
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90.847 Million cell updates/sec
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                                                                      January 26, 2005, 15:54:02; Search time 14.6 Seconds
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score greater than or equal to the score of the result bein
and is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
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US-09-142-524D-44

US-09-142-524D-44

US-09-142-524D-42

US-08-166-568E-91

US-08-469-260A-366

US-08-469-260A-366

US-08-445-50A-366

US-08-424-550B-366

US-09-445-524D-45

US-09-445-524D-45

US-09-440-884A-248

US-09-46-421B-248

US-09-46-421B-248

US-09-630-940B-248

US-09-247-527-18

US-09-308-368-13

US-09-308-368-13
                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-955-138-115
US-09-252-586-27
                                                                                                                                                                                                        478139 seqs, 66318000 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                             OM protein - protein search, using sw model
                                                                                                                                104
1 GVVPVHAQDGDAITMRNVTD 20
                                                                                                                                                                    BLOSUM62 Gapext 0.5
                                                                                                                                                                                                                                                                                                                                             Issued Patents AA:*
                                                                                                                      US-09-202-464-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query
Match Length
        Copyright
                                                                                                                                                                                                                                                        Minimum DB seq length: 0 Maximum DB seq length: 20
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                                                                                                                                Perfect score:
                                                                                                                                                                     Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Database :
                                                                                                                                                Sequence:
                                                                                                                                                                                                         Searched:
                                                                      Run on:
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No.
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28 26.9 19 4 US-09-512-563C-43 Sequence 43, Appl 29 28 26.9 20 4 US-08-845-381E-48 Sequence 43, Appl 31 27 26.0 16 2 US-08-612-402B-3 Sequence 22, Appl 32 27 26.0 16 2 US-08-490-473B-17 Sequence 27, Appl 34 27 26.0 16 3 US-08-915-213-17 Sequence 17, Appl 35 27 26.0 16 3 US-08-915-213-17 Sequence 17, Appl 36 27 26.0 18 3 US-09-255-217-17 Sequence 17, Appl 36 27 26.0 18 3 US-09-31-17-17 Sequence 17, Appl 37 27 26.0 18 3 US-09-31-17-17 Sequence 17, Appl 37 27 26.0 18 3 US-09-31-17-17 Sequence 2, Appl 37 27 26.0 18 3 US-09-31-27-218-5 Sequence 2, Appl 40 26.5 25.5 17 3 US-08-215-721B-5 Sequence 36, Appl 41 26.5 25.5 17 3 US-08-215-035-8 Sequence 36, Appl 42 26.5 25.0 11 3 US-08-215-805A-2 Sequence 36, Appl 44 26 25.0 11 3 US-09-48-338-15 Sequence 15, Appl 44 26 25.0 11 4 US-09-88-838A-15 Sequence 15, Appl 45 26 25.0 11 4 US-09-88-838A-15 Sequence 1, Appl 16-09-89-15-17-17 Sequence 15, Appl 45 26 25.0 12 2 US-08-487-675-1
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## ALIGNMENTS

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US-08-467-023-40

Sequence 40, Application US/08467023

Patent No. 600386

GENERAL INPORMATION:

APPLICANT: Griffeeh, Irwin J.;

APPLICANT: Garman, Aichard D;

APPLICANT: Garman, Aichard D;

APPLICANT: Boulock, Joanne;

APPLICANT: Bauer, Andrew;

APPLICANT: Exley, Mark P;

APPLICANT: Brauer, Andrew;

APPLICANT: Brauer, Andrew;

APPLICANT: Dowers, Steven P;

ITILE OF INVENTION: Japanese Cedar Pollen

NUMBER: OF SEQUENCES: 261

CORRESPONDENCE ADDRESS: 3

ADDRESSEE: 1nmulogic Pharmaceutical Corporation, Inc.

STREET: 610 Lincoln St

CONFORTE: 1NP PC COMPARIED PORM:

MEDIU TYPE: Floppy disk

COMPUTER: INP PC COMPARIED PORM:

MEDIU TYPE: Floppy disk

COMPUTER: INP PC COMPARIED PORM:

MEDIU TYPE: Hoppy disk

COMPUTER: INPRAFITION NUMBER: US/08/467, 023

FILING DATE: June 6, 1995

CLASSIFICATION NUMBER: 08/350, 225

FILING DATE: December 6, 1994

ATTORNEY AGENT INPORMATION:

TELEPHONE: (617) 227 7400

TELEPHONE: (617) 227 7400

TELEPHONE: (617) 227 7594

INTORMATION FOR SEQ ID NO: 40:

SEGUENCE CHARACTERISTICS:

LEMENTH: 20 amino acids

TYPE: TYPE: Inhear APPLICATION

NUMBER: 20 amino acids

TYPE: MARKEN TYPE: Inhearnal
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APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
FURENT FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID 0.42
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APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bonlock, Jonene;
APPLICANT: Bond, Unian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Youn, Mei-Chang;
APPLICANT: Youn, Mei-Chang;
APPLICANT: Young, Slui-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Brauer, Andrew;
APPLICANT: Bowers, Steven P.;
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 15;
                                                                Length 15;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (1). (15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 28
US-09-142-524D-42
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 30 US-09-142-524D-44
                                                                DB 4;
0.008;
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Pred. No. 0.61;
0; Mismatches
                                                                                                              3; Mismatches
                                                                51.9%; Score 54; 60.0%; Pred. No.
                                                                                                                                                                                                                                                                                                         Sequence 42, Application US/09142524D Patent No. 6719976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 39, Application US/08467023 ; Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 80.0%;
Matches 8; Conservative
                                                                                                                                                        6 HAQDGDAITMRNVTD 20
                                                                                                                                                                                    610 Lincoln St
                                                                                   Best Local Similarity 60.0
Matches 9, Conservative
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                                                                                                                                                                                                                                                                                                                                     Patent No. 6719976
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Waltham
                                                                                                                                                                                                                                                                                         US-09-142-524D-42
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                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE REPERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dairik, Kazuo
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Kabsuke
APPLICANT: Kino, Kabsuke
APPLICANT: Kino, Kabsuke
APPLICANT: No, Kabsuke
FILE REFERRNCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 44
LENGTH: 15
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                                         Score 73; DB 3; Length 20;
Pred. No. 6.3e-06;
3; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KAX: MISC_FEATURE
LOCATION: (1)...(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 29
US-09-142-524D-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-142-524D-44; Sequence 44, Application US/09142524D; Patent No. 6719976
                                                                                                                                                                                                                                                                                     Sequence 43, Application US/09142524D
Patent No. 6719976
                                                                                                                                   1 GVVPVHAQDGDAITMRNVTD 20
                                                                                                                                                               1 GVEPVHPQDGDALTLRTATN 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Cryptomeria japonica
                                              70.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GVVPVHAQDGDAITM 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 65.09
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(15)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       US-09-142-524D-43
    US-08-467-023-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 43
LENGTH: 15
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APPLICANT: ISA K. MUSTAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4; Length 16;
51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                  STATE: LEGIT FRAN.
STATE: LL
CONNTRY: USA
ZIP: 60064-3500
COMPOTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,260A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                             D377/AP6D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.8%; Score 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5527.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
                                                      JOHN N. SIMONS
TAMI J. PILOT-MATIAS
GEORGE J. DAMSON
GEORGE G. SCHLAUDER
SURESH M. DESAI
THOWAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BUIJK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JOHN N. SIMONS
TAMI J. PILOT-MATIAS
GEORGE J. DAWSON
GEORGE G. SCHLAUDER
SURESH M. DESAI
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BUIJK
  Application US/08469260A
                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: PORREMENT, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.F
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 366:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 366, Application US/08488446; Patent No. 6558898 GENERAL INFORMATION: GENERAL TAPONN N. SIMONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
XGY: linear
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US-08-488-446-366
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APPLICANT:
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APPLICANT:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: Bhattacharjee, J.

APPLICANT: Bhattacharjee, J.

APPLICANT: Bhattacharjee, Vasker

APPLICANT: Bhattacharjee, Vasker

TITLE OF INVENTION: METHODS AND REAGENTS FOR DETECTING FUNGAL PATHOGENS IN

TITLE OF INVENTION: A BIOLOGICAL SAMPLE

FILE REFERENCE: 96,247-A

CURRENT APPLICATION NUMBER: US/09/106,568E

CURRENT FILING DATE: 1998-06-29

PRIOR PILING DATE: 1997-05-20

NUMBER OF SEQ ID NOS: 160

SOFTWARE: Microsoft Word 97

LENGTH: 19
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CTHER INFORMATION: Polypeptide segment of ACVS_CEPAC shown in Figure 4. US-09-106-568E-91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               35.6%; Score 37; DB 4; Length 19; 66.7%; Pred. No. 8.7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                            OPERATING SYSTEM: PC-LUCS, FIG.
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILLING DATE: June 6, 1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
REGISTRALION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7400
TELEPRAM: (617) 227-5941
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 6
US-09-106-568E-91
; Sequence 91, Application US/09106568E
; Patent No. 6455248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 80.0
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 66.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GVVPVHAQDG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11 GVEPVHPQDG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-467-023-39
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Gaps

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US-08-469-260A-366

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: LENGTH: 16 amino acids
; TOPE: amino acid
; TOPOLCY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 366:
US-08-467-344A-366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: ABBOTT LABORATORIES STREET: 100 ABBOTT PARK ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION
TELEPANCE: 708-937-6365
TELEPAK: 708-938-2623
INFORMATION FOR SEQ ID NO: 366:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435435.
ATTORNAY AGENT INFORMATION:
NAME: POREMBEKI, PRIGISILLA E.
REGISTRATION 'NUMBER: 33,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.4
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 VPVHAQ 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 VPVHAQ 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 10
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US-08-467.344A-366
US-08-467.34A-366
US
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APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT LABORATORIES D377/AP6D
CITY: ABBOTT PARK
STATE: 11
COUNTY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: 
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51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REPERENCE/DOCKET NUMBER: 5527.PC.01
REPERENCE/DOCKET NUMBER: 5527.PC.01
REPERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEFAX: 708-938-2635
INFORMATION FOR SEQ ID NO: 366:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
STATE: 1L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.8%; Score 32; 100.0%; Pred. No.
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ZIP: 60064-3500
COMPUTER READABLE FORM:
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Matches 6; Conservative
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MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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Sequence 366, Application US/08424550B

Sequence 366, Application US/08424550B

Patent No. 6720166

GENERAL INFORMATION:
APPLICANT: TAMI JOHN N. SIMONS
APPLICANT: TAMI JOHN SIMONS
APPLICANT: TAMI JOHN SIMONS
APPLICANT: TAMI SECREG J. DAWSON
APPLICANT: SURESH M. DESAI
APPLICANT: SURESH M. DESAI
APPLICANT: SURESH M. DESAI
APPLICANT: SHERI L. BUIJK
APPLICAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,344A
FILING DATE: 07-Jun-1995
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/424,550
FILING DATE: <Unknown>
ATTORNEY/AGENT INPORMATION:
NAMM: POREMBSKI, PRISCILLA E.
REDISTANION NUMBER: 33,207
REPERENCE/DOCKET NUMBER: 5527.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
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CONTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              D377/AP6D
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APPLICANT: Henderson, Robert A.
APPLICANT: McNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C11
CURRENT APPLICATION NUMBER: US/09/643,597
CURRENT FILING DATE: 2000-08-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28.8%; Score 30; DB 3; Length 20; 50.0%; Pred. No. 1.5e+02;
                                                                 ImmuLogic Pharmaceutical Corporation, Inc.
                                                                                                                                                                                                                                                                                                                                               FILING DATE: June 6, 1995
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7401
                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
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1.5e+02;
2;
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Patent No. 6426072
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Hosken, Nancy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Skeiky, Yasir A.W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kalos, Michael D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGIH: 20 amino acids
                                                                                        STREET: 610 Lincoln St
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Li, Samuel X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5; Conservative
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Fan, Liqun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                 CORRESPONDENCE ADDRESS: ADDRESSE: ImmuLogic
                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 DAITMRNVTD 20
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ORGANISM: Homo sapiens
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SOFTWARE: FastSEQ for
    LITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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Best Local Similarity
                                                                                                                 Waltham
                                                                                                                                                            USA
                                                                                                                                                                            ZIP: 02154
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                                                                                                                                     STATE: M
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-467-023-41
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LENGTH: 20
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Kazue, Akinori
APPLICANT: Iwama, Akiko
FILE REFERENCE: SPO-103
CURRENT PAPLICATION WUMBER: 1988-09-09
PRIOR APPLICATION WUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
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APPLICANT: Griffeth, Joanne;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Graman, Richard D;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Braver, Andrew;
APPLICANT: Exper, Mark A.;
APPLICANT: Expers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28.8%; Score 30; DB 4; Length 15; 50.0%; Pred. No. 1e+02; ative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                DB 4; Length 16; 51;
                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: MISC_FEATURE
COCATION: (1)...(15)
COCHER INFORMATION: Cryjl peptide, Figure 1, Row 31
MS-09-142-524D-45
                                                                                                                                                                                                                                                30.8%; Scor.
100.0%; Pred. No. ...
    5527.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 45, Application US/09142524D Patent No. 6719976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41, Application US/08467023
Patent No. 6090386
                                                          TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 366:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 aming acids
                      TELECOMMUNICATION INFORMATION TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 174.
SOFTWARE: Patentin version 3.1
SEQ ID NO 45
LENGTH: 15
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
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Best Local Similarity 50.0
Matches 5; Conservative
                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: protein
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Gaps

2 ETGDPVTLRLLDD 14

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Search completed: January 26, 2005, 16:08:24 Job time : 15.6 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 20;
                                         28.8%; Score 30; DB 4; Length 20; 38.5%; Pred. No. 1.5e+02;
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Patent No. 6518256
GENERAL INFORMATION:
APPLICANT: Fan, Liqun
APPLICANT: Fanger, Anchael D.
APPLICANT: Hosken, Nancy A.
APPLICANT: Fanger, Cary R.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
TITLE REFERENCE: 210121.455C8
CURRENT APPLICATION NUMBER: US/09/542,615A
CURRENT FILLING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 248
LENTH: 20
                                                                                                                                                                                                                                                                    Sequence 248, Application US/09480884A
Fatent No. 6482597
GENERAL INFORMATION:
APPLICANT: Wang, Tongtong
APPLICANT: Fan, Liqun
APPLICANT: Fan, Liqun
APPLICANT: Fan, Liqun
APPLICANT: Fanger, Nancy A.
APPLICANT: Fanger, Michael D.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C6
CURRENT FILING DATE: 2001-08-27
NUMBER OF SEQ ID NOS: 330
SOFTWARE: FaatsEQ for Windows Version 3.0
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 30; DB 4;
Pred. No. 1.5e+02;
4; Mismatches 4
                                                                                    4; Mismatches
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Best Local Similarity 38.5%;
Matches 5; Conservative '
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2 ETGDPVTLRLLDD 14
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Best Local Similarity 38.55
Matches 5; Conservative
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2 ETGDPVTLRLLDD 14
                                      Query Match
Best Local Similarity 38.5
Matches 5; Conservative
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CRGANISM: Homo sapiens
US-09-542-615A-248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT; ORGANISM: Homo sapiens
US-09-480-884A-248
                                                                                                                                                                                                                                      RESULT 14
US-09-480-884A-248
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US-09-542-615A-248
US-09-643-597-248
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8 QDGDAITMRNVTD 20

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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1608061 seqs, 361289386 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Published_Applications_AA:*
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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107
1 DAITMRNVTDVWIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Disea

Description	Sequence 45, Appl	Sequence 46, Appl		~	Sequence 211, App	Sequence 41072, A	Sequence 32, Appl	Sequence 32, Appl	Sequence 33, Appl	Sequence 30, Appl	Sequence 30, Appl	Sequence 30, Appl	Sequence 33, Appl
SUMMARIES	US-10-354-240-45	US-10-354-240-46	US-10-354-240-47	US-10-225-567A-1876	US-10-473-127-211	US-09-864-761-41072	US-10-011-095-32	US-10-010-667A-32	US-10-165-044-33	US-10-408-009-30	US-10-856-109-30	US-10-857-785-30	US-10-753-195-33
DB	14	14	14	14	17	6	14	14	14	15	17	17	17
% Query Match Length DB	15	15	15	16	13	19	15	15	15	15	15	15	15
* Query Match	57.9	54.2	43.9	37.4	34.6	33.6	32.7	32.7	32.7	32.7	32.7	32.7	32.7
Score	62	28	47	40	37	36	35	35	35	35	35	35	35
Result No.	-	8	e	4	ß	9	7	80	σ	10	11	12	13

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Gaps

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Query Match 57.9%; Score 62; DB 14; Length 15; Best Local Similarity 60.0%; Pred. No. 0.007; Matches 9; Conservative 4; Mismatches 2; Indels

1 DAITMRNVTDVWIDH 15

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Sequence 88, Appl Sequence 1795, Ap Sequence 1745, Ap Sequence 92, Appl Sequence 235, App Sequence 235, App Sequence 235, App Sequence 6, Appl Sequence 6, Appl Sequence 46, Appl Sequence 42, App Sequence 420, App Sequence 420, App Sequence 421, App Sequence 421, App	Sequence 440, App Sequence 105, App Sequence 105, App Sequence 105, App Sequence 405, App Sequence 405, App Sequence 56, App Sequence 64, Appl Sequence 44, Appl Sequence 198, App Sequence 198, App Sequence 198, App Sequence 198, App Sequence 198, App Sequence 198, App	Agent for Treating Allergic	
US-10-432-422-88 US-10-225-567A-1795 US-09-995-529-92 US-09-995-529-92 US-09-995-529-92 US-09-995-529-92 US-10-653-595-335 US-10-63-3595-335 US-10-1312-311-6 US-10-1312-311-6 US-10-084-813-46 US-09-793-451-424 US-09-793-451-424 US-09-793-451-424 US-10-283-722-440 US-10-283-722-440	US-10-283-903-440 US-10-283-903-440 US-09-793-451-105 US-00-793-451-105 US-10-283-722-105 US-10-283-903-105 US-10-283-903-105 US-10-285-6493-4 US-10-285-6493-4 US-10-378-173-45 US-10-378-173-45 US-10-609-217-198 US-10-609-217-198 US-10-612-388-198 US-10-645-761-198	54240. sed Immunotherapeutic 29 J997/00740 9/142,524 a	c mov /1 pin613
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115 4 116 117 118 118 119 118 118 118 118 118 118 118	<b>๛</b>	RESULT 1  US-10-354-240-45  Sequence 45, Applicatio Publication No. US20030 GENERAL INFORMATION: APPLICANT: Kume, Akin FULEN ERPERENCE: SPO-10 CURRENT APPLICATION NUMB PRIOR FILING DATE: 199 NUMBER OF SEQ ID NOS: SOFTWARE: PATENTIN OF SEQ ID NOS: CROTHARE: PATENTIN OF SEQ ID NOS: CROTHARE: PATENTIN OF SEQ ID NOS: CROTHARE: MASC FEATUR CROTHER: NAME/KEY: MISC FEATUR COTHER INFORMATION: CTTER	US-10-354-240-45

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APPLICANT: LifeSpan Biosciences
APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARES: Patentin version 3.1
SEQ ID NO 1876
LENGTH: 16
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                           ;
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TITLE OF INVENTION: TRANSLATIONAL PROFILING FILE REFERENCE: 0819-1026W01.

CURRENT PEDLICATION NUMBER: 05/279,495

PRIOR APPLICATION NUMBER: 60/279,495

PRIOR FILING DATE: 2001-03-28

PRIOR FILING DATE: 2001-05-21.

PRIOR PILING DATE: 2001-05-21.

PRIOR PILING DATE: 2001-06-08

PRIOR FILING DATE: 2001-09-08

PRIOR FILING DATE: 2001-10-01.

PRIOR FILING DATE: 2001-10-04.

PRIOR FILING DATE: 2001-12-04.

PRIOR FILING DATE: 2001-12-04.
  Pred. No. 1.6;
2; Mismatches
                                                                                                                                                                                                                                                             Sequence 1876, Application US/10225567A Publication No. US20030113798A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 211, Application US/10473127; Publication No. US20040236091A1; GENERAL INFORMATION:
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70.0%;
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  Best Local Similarity 70.0
Matches 7; Conservative
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Best Local Similarity 54.5
Matches 6; Conservative
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1 IWIDHNSFSN 10
                                                                               11 VWIDHNSLSD 20
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ORGANISM: Homo sapiens
US-10-473-127-211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
US-10-225-567A-1876
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                   APPLICANT: Sone, Toshio:
APPLICANT: Sone, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Ivama, Akinori
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Akiko
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT APPLICATION NUMBER: US/10/354,240
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 46
LENGTH: 15
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT APPLICATION NUMBER: US/10/354,240
PRIOR FILING DATE: 2003-01-29
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 47
TYPE: RIT
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Pred. No. 0.03;
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COCATION: (1)...(15)

OTHER INFORMATION: Cryjl peptide, Figure 1, Row 33

US-10-354-470-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: MISC_FEATURE
CCATION: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 32
US-10-354-240-46
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                                                                                                                                 Sequence 46, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Cryptomeria japonica
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 60.0%;
Matches 9; Conservative
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        1 DALTLRTATNIWIDH 15
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                                                                                                                 US-10-354-240-46
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APPLICANT: Afar Daniel
APPLICANT: Hubert, Rene S.
APPLICANT: Hubert, Rene S.
APPLICANT: Leong, Kahan
APPLICANT: Astiano, Arthur B.
APPLICANT: Raitano, Douglas C.
APPLICANT: Mitchell, Steve Chappell
TITLE OF INVENTION: ANTHODIES IMMUNOSPECIFIC FOR STEAPI (AS AMENDED)
FILE REPRENCE: 511582001610
CURRENT FILING DATE: 1050-12-06
FRIOR APPLICATION NUMBER: 09/323,873
PRIOR PELING DATE: 1999-06-01
PRIOR PELING DATE: 1999-06-01
PRIOR PELING DATE: 1999-06-30
NUMBER OF SEQ ID NOS: 32
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 32
LENTH: 15
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; Sequence 32, Application US/10010667A
; Publication No. US20030055217A1
; GENERAL INFORMATION:
    APPLICANT: Atar, Daniel
; APPLICANT: Hubert, Rene S.
; APPLICANT: Leong, Kahan
; APPLICANT: Ratiano, Arthur B.
; APPLICANT: Saffran, Douglas C.
; APPLICANT: Mitchell, Steve Chappell
; TILE OF INVENTION: NOVEL SERRENTINE TRANSMEMBRANE ANTIGENS
; TILE OF INVENTION: BEYRESSED IN HUMAN CANCERS AND USES THEREOI
; FILE REFERENCE: 511582001601
; CURRENT APPLICATION NUMBER: US/10/10,667A
; CURRENT APPLICATION NUMBER: 60/087,520
; PRIOR FILING DATE: 1999-06-01
; PRIOR FILING DATE: 1998-06-01
; PRIOR APPLICATION NUMBER: 60/091,183
; PRIOR APPLICATION NUMBER: 60/091,183
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                                    Length 19;
                                                                                              3; Indels
                              Score 36; DB 9; Pred. No. 1.1e+02; 3; Mismatches 3
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                            US-10-011-095-32
; Sequence 32, Application US/10011095
; Publication No. US20030045682A1
; GENERAL INFORMATION:
                                    33.6%;
                              Query Match
Best Local Similarity 50.0
Matches 6; Conservative
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Best Local Similarity 50.0
Matches 5; Conservative
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CORGANISM: Homo sapiens
US-10-011-095-32
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                                                                                                                                                                             Sequence 41072, Application US/09864761
Patent No. US200200487531
GENERAL INFORMATION:
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Chen, Wesheng
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aconica-X-1
CURRENT APPLICATTON NUMBER: US/09/864,761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EXPRESSED IN BONE MARROW, SIGNAL = 3
EXPRESSED IN FETAL LIVER, SIGNAL = 3.5
EST_HOMAN HIT: A1719171.1, EVALUE 4.00e-04
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EXPRESSED IN PLACENTA, SIGNAL = 2.6
EXPRESSED IN ADULT LIVER, SIGNAL = 3.7
EXPRESSED IN HEART, SIGNAL = 2.6
EXPRESSED IN BRAIN, SIGNAL = 2.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR PILING DATE: 2000-02-04
PRIOR PILING DATE: 2000-02-05
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/236,356
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR PILING DATE: 2001-01-30
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1 VFLDHNALPD 10
   11 VWIDHNSLSD 20
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ORGANISM: Homo sapiens
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OTHER INFORMATION:
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FILE REFERENCE: 51582016015

FILE REFERENCE: 51582016016

CURRENT APPLICATION NUMBER: US/10/408,009

CURRENT FILING DATE: 2003-04-04

FRIOR PEPLING DATE: 2094-05,486

PRIOR FILING DATE: 1999-12-06/15

PRIOR FILING DATE: 1999-12-06-01

NUMBER OF SEQ ID NOS: 36

SOFTWARE: FASLESQ for Windows Version 4.0

LENGTH: 15

LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rene S. Hubert
APPLICANT: Rene S. Hubert
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Stephen C. Mitchell
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
TITLE OF INVENTION: BENEBESED IN HUMAN CANCERS AND USES THEREOF
FILE REFERENCE: 511582001606
CURRENT APPLICATION NUMBER: US 09/455,486
FRIOR APPLICATION NUMBER: US 09/455,486
PRIOR FILING DATE: 1999-106-01
PRIOR FILING DATE: 1999-06-01
PRIOR FILING DATE: 1999-06-01
PRIOR FILING DATE: 1999-06-01
PRIOR FILING DATE: 1999-06-01
PRIOR FILING DATE: 1998-06-01
SPRIOR FILING DATE: 1998-06-01
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Pred. No. 1.2e+02;
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; Sequence 30, Application US/10857785
; Publication No. US20040219162A1
; GENERAL INFORMATION:
; APPLICANT: Daniel B. Afar
; APPLICANT: Rene S. Hübert
; APPLICANT: Arthur B. Raitano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 30, Application US/10856109 Publication No. US20040219591A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      32.7%;
50.0%;
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50.0%;
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Best Local Similarity 50...
Best Local Si Conservative
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Best Local Similarity 50.0
Matches 5; Conservative
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CRGANISM: Homo sapiens
US-10-408-009-30
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APPLICANT: Agensy, Inc.
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
ITITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
ITITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
ITITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
ITITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
ITITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
ITITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
ITITLE OF INVENTION NUMBER: US 60/081,183
PRIOR FILING DATE: 1998-06-01
PRIOR PELING DATE: 1998-06-30
PRIOR PELING DATE: 1999-06-30
PRIOR PELING DATE: 1999-06-30
PRIOR PELING DATE: 1999-06-01
PRIOR PLING DATE: 1999-06-01
PRIOR PLING DATE: 1999-06-01
PRIOR PLING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 57
SOFTWARE: FastSEQ for Windows Version 4.0
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Publication No. US20040072196A1
GENERAL INFORMATION:
APPLICANT: Daniel E. Afar
APPLICANT: Rene S. Hubert
APPLICANT: Arthur B. Raitano
APPLICANT: To Douglas C. Saffran
APPLICANT: Stephen C. Mitchell
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; Indels
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Pred. No. 1.2e+02;
2; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 33, Application US/10165044 Publication No. US20030149531A1 GENERAL INFORMATION:
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50.0%;
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Best Local Similarity 50.0
Matches 5; Conservative
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                                                            SEQ ID NO 32
LENGTH: 15
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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Best Local Similarity
Matches 5; Conserva
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US-10-408-009-30
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US-10-165-044-33
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Gaps

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APPLICANT: LifeSpan Biosciences
APPLICANT: Enown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Janna C.
APPLICANT: Burmer, Janna C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS FILE REFERENCE: 1920-4-4
CURRENT APPLICATION NUMBER: 2001-12-19
PRIOR PILING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
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                            Indels
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Syngenta Participations AG
APPLICANT: Cornell Research Foundation, Inc.
APPLICANT: Toder, Olen
APPLICANT: Turgeon, Barbara G.
APPLICANT: Turgeon, Barbara G.
APPLICANT: Lu, Shen-wen
TITLE OF INVENTION: Fungal Iron Reductase Gene
FILE REFERENCE: 1360.017W01
CURRENT APPLICATION NUMBER: US/10/432,422
CURRENT FILING DATE: 2003-05-21
PRIOR APPLICATION NUMBER: US 60/252,732
PRIOR FILING DATE: 2000-11-22
PRIOR FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 210
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 34; DB 15;
Pred. No. 1.6e+02;
3; Mismatches 2
  50.0%; Pred. No. 1.2e+02;
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                            2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 1795, Application US/10225567A; Publication No. US20030113798A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   i LENGTH: 14
; TYPE: PRT
; ORGANISM: Cochliobolus heterostrophus
US-10-432-422-88
                                                                                                                                                                                                                                                                     Sequence 88, Application US/10432422
Publication No. US20040076981A1
GENERAL INFORMATION:
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50.0%;
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SOFTWARE: Patentin version 3.1
SEQ ID NO 1795
LENGTH: 20
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Best Local Similarity 50.0
Matches 5; Conservative
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                         5; Conservative
                                                                               6 RNVTDVWIDH 15
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6 QNKEDAWIEH 15
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2 EIWVDSPSLS 11
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US-10-225-567A-1795
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APPLICANT: Agensy, Inc.
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Arthur B. H. Afar
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: BERPENTINE TRANSMEMBRANE ANTIGENS
TITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
FILE REFERENCE: 5158-2004-01-06
FILE REFERENCE: 5158-2004-01-06
PRIOR APPLICATION NUMBER: US 60/091,520
PRIOR APPLICATION NUMBER: US 60/091,183
PRIOR PLING DATE: 1998-06-01
PRIOR PLING DATE: 1998-06-01
PRIOR APPLICATION NUMBER: US 09/455,486
PRIOR APPLICATION NUMBER: US 09/455,486
PRIOR PLING DATE: 1999-06-01
PRIOR PLING DATE: 2000-12-06
PRIOR PLING DATE: 2000-12-06
PRIOR PLING DATE: 2000-12-06
PRIOR PLING DATE: 2000-12-06
PRIOR PRIOR PRING DATE: 2000-13040
PRIOR PLING DATE: 2000-12-06
PRIOR PRING DATE: 1999-06-01
PRIOR PLING DATE: 1999-06-01
                 APPLICANT: Stephen C. Mitchell
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE AND USES THEREOF
FILE REFERENCE: 5014-01607
CURRENT APPLICATION NUMBER: US/10/857,785
CURRENT FILING DATE: 2004-05-28
PRIOR APPLICATION NUMBER: US 09/455,486
PRIOR PILING DATE: 1999-12-06
PRIOR APPLICATION NUMBER: US 60/087,520
PRIOR APPLICATION NUMBER: US 60/091,183
PRIOR PILING DATE: 1998-06-01
PRIOR APPLICATION NUMBER: US 60/091,183
PRIOR PILING DATE: 1998-06-01
PRIOR APPLICATION NUMBER: US 60/091,183
PRIOR PILING DATE: 1998-06-01
PRIOR APPLICATION NUMBER: US 60/091,183
PRIOR PILING DATE: 1998-06-01
SPRIOR PILING DATE: 1998-06-01
SPRIOR PILING DATE: 1998-06-01
SPRIOR PILING DATE: 1998-06-01
SPRIOR APPLICATION NUMBER: US 60/091,183
PRIOR APPLICATION NUMBER: US 60/091,183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 35; DB 17;
Pred. No. 1.2e+02;
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Publication No. US20050004349A1
GENERAL INFORMATION:
  Douglas C. Saffran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    32.7%;
50.0%;
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Best Local Similarity 50.0
Matches 5; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Search completed: January 26, 2005, 16:54:58 Job time : 56.5 secs

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90.847 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
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US-08-467-023-212

US-08-467-023-211

US-09-142-524D-47

US-08-67-023-42

US-08-23-873A-32

US-08-981-122-77
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US-08-370-567-5
US-08-438-759-5
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US-08-466-033-85
US-08-466-033-89
US-08-444-733-85
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US-08-640-877-22
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Maximum Match 100%
Listing first 45 summaries
                                                           OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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107
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Match Length DB
           Copyright
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Perfect score:
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Sequence 12, Appl	US-08-818-253-12	7	17	26.2	28	45
	PCT-US92-10068-14	w.	16	26.2	58	44
Seguence 30, Appl	US-09-914-695-30	4	16	26.2	28	43
Sequence 12, Appl	US-08-318-200-12	Н	16	26.2	58	42
Seguence 59, Appl	US-09-069-827A-59	4,	12	26.2	28	41
Sequence 3, Appli	US-08-135-319A-3	4	10	26.2	28	. 04
Sequence 2, Appli	US-08-135-319A-2	4	10	26.2	28	39
Sequence 7, Appli	US-08-430-925A-7	ч	80	26.2	28	38
Sequence 6, Appli	US-08-787-547-6	Н	7	26.2	58	37
Sequence 73, Appl	PCT-US95-06266-73	ഗ	50	27.1	53	36
Seguence 69, Appl	PCT-US95-06266-69	ហ	20	27.1	53	35
Sequence 89, Appl	US-08-485-910-89	~	20	27.1	59	34
Sequence 85, Appl	US-08-485-910-85	~	20	27.1	53	33
_	US-08-461-361-89	~	20	27.1	59	32
	US-08-461-361-85	~	20	27.1	53	31
Sequence 89, Appl	US-08-464-134-89	7	20	27.1	53	30
Sequence 85, Appl	US-08-464-134-85	~	20	27.1	29	29
Sequence 89, Appl	US-08-444-733-89	~	20	27.1	59	28

## ALIGNMENTS

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Patent No. 6090366

Patent No. 6090366

GENERAL INPORMATION:

APPLICANT: Garma, Richard D;

APPLICANT: Garma, Richard D;

APPLICANT: Pollock, Joanne,

APPLICANT: Band, Julian F;

APPLICANT: Bond, Julian F;

APPLICANT: Brauer, Andrew;

ADDRESSED SECONTES: 261

CONTRX: 04

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen TITLE OF SEQUENCES: 261
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
                                                                                                          Length 15;
                                                                                                                                                                       3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION UNMBER: 08/350,225
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTOMERY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE / DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
, OTHER INFORMATION: Cryjl peptide, Figure 1, Row 32
US-09-142-524D-46
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Pred. No. 0.0097;
                                                                                                             Score 58; DB 4;
Pred. No. 0.0069;
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Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Brauer, Andrew;
APPLICANT: POWERS, Steven P.
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 amino acids
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; FRAGMENT TYPE: internal
US-08-467-023-203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                   Query Match
Best Local Similarity
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Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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APPLICANT: Ivana, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REPRENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 45
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFRENCE: SPO-103
CURRENT PELLING DATE: 1990-09-09
CURRENT FILING DATE: 1990-09-09
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 46
LEASTH NOS: 174
LEASTH NOS:
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                                                                                72.0%; Score 77; DB 3; Length 20; 60.0%; Pred. No. 1e-05; 1ve 5; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (1) ...(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 31
US-09-142-524D-45
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 45, Application US/09142524D
Patent No. 6719976
GENERAL INFORMATION:
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                                                                                                                                                                                                                 1 DAITMRNVTDVWIDHNSLSD 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DAITMRNVTDVWIDH 15
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1 DALTLRTATNIWIDH 15
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APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
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Best Local Similarity 60.0
Matches 9; Conservative
                                                                                                                Best Local Similarity 60.0 Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: MISC_FEATURE
                                                                                                                                                                                                                                                                                                                                                                     RESULT 2
US-09-142-524D-45
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                             US-08-467-023-41
                                                                                       Query Match
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APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From
TITLE OF INVENTION: Japanese Cedar Pollen
TITLE OF SEQUENCES:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
STATE: MA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Algenese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 53; DB 3; Length 16;
Pred. No. 0.046;
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                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEFRAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 212:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: AMINO acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 211, Application US/08467023
; Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 66.7%;
Matches 8; Conservative 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 RNVTDVWIDHNS 17
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RTATNIWIDHNS 13
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
US-08-467-023-211
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APPLICANT: Griffeet, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Applicant Application, Inc.
STREET: MA
COUNTRY: WSA
ZIP: A...
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COMPUTER: FLODPY OLSK
COMPUTER: PATENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION ATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-740
TELEFRA: (617) 227-740
TELEFRA: (617) 227-5941
INFORMATION FOR SEQ ID NO: 210:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
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Patent No. 6090386
GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bollock, Joanne;
APPLICANT: Bond, Julian F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
Garman, Richard D;
Kuo, Mei-Chang;
Yeung, Siu-mei H.;
Brauer, Andrew;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51.4%;
60.0%;
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Best Local Similarity 60.0
Matches 9; Conservative
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; FRAGMENT TYPE: internal
US-08-467-023-210
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COMPUTER READABLE FORM:
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APPLICANT:
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APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Brauer, Andrew;
APPLICANT: Every, Mark A.;
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
ADDRESSEE: ADDRESSE: ADDRESSE: ADDRESSEE: ADDRESSEE: ADDRESSEE: ADDRESSEE: STREET: 610 Lincoln St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 43.9%; Score 47; DB 3; Length 20; Best Local Similarity 70.0%; Pred. No. 0.52; Matches 7; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATION STEED PC-DOSING-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (617) 227-7400
TELEFAX: (617) 227-7401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 32. Application US/09323873A patent No. 6329503 GENERAL INFORMATION: APPLICANT: Daniel E. Afar APPLICANT: Rene S. Hubert APPLICANT: Rahan Leong APPLICANT: Arthur B. Raitano APPLICANT: Arthur B. Raitano APPLICANT: Steve Chappell Mitchell
                                                                   Sequence 42, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
                                                                                                                                         Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
Garman, Richard D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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MEDIUM TYPE: Floppy
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CITY: Waltham
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ZIP: 02154
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                                                  JS-08-467-023-42
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APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Mama, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwana, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REPRENCE: SPO-103
CURRENT PILING DATE: 1998-09-09
PRIOR PLING DATE: 1997-00-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 47
LENGTH: 15
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                                                                                                                                                                                      CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 211:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 mming acids
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OFERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: June 6, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (1).7(15)
CTHER INFORMATION: Cryjl peptide, Figure 1, Row 33
US-09-142-524D-47
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ORGANISM: Cryptomeria japonica
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Best Local Similarity 70.0
Matches 7: Conservative
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Best Local Similarity 66.7
Matches 8; Conservative
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4 RTATNIWIDHNS 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : | | | | | | :
1 IWIDHNSFSN 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
GY: linear
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TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
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Patent No. 5656726
GENERAL INFORMATION:
APPLICANT: Rosenberg, Steven
APPLICANT: Doyle, Michael
APPLICANT: Goodson, Robert
TITLE OF INVENTION: Peptide Inhibitors of Urokinase Receptor TITLE OF INVENTION: Activity
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28.0%; Score 30; DB 4; Length 15; 50.0%; Pred. No. 1.7e+02; tive 3; Mismatches 2; Indel8
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/370,567
                                              FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
FRICH APPLICATION NUMBER: PCT/JP97/00740
FRICH APPLICATION NUMBER: PCT/JP97/00740
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
LENGTH: 15
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CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/061,514
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ATTORNEY/AGENT INFORMATION:
NAME: Green, Grant D.
REGISTRATION NUMBER: 31,259
REFRENCE/POCKET NUMBER: 0941
TELECOMMUNICATION INFORMATION:
TELEPHONE: 510-601-2706
                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 4560 Horton Street
Emeryville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 28.0
Best Local Similarity 50.0
Matches 5; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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OTHER INFORMATION: Sequence of a peptide synthesized in Comparative Example 2 from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatiomi, Masaharu
ITILE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: DT/JP96/01734
PRIOR PILING DATE: 1996-06-21
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR PILING DATE: 1996-06-21
SOFTWARE: Patentin Ver. 2.0
SSQ ID NO 77
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; TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
; TITLE OF INVENTION: EXPRESSED IN HUMAN CANCERS AND USES THEREOF
; FILE REFERENCE: 129.16USU2
; CURRENT APPLICATION NUMBER: US/09/323,873A
; CURRENT FILING DATE: 1999-06-01
; PRIOR APPLICATION NUMBER: 60/091,183
; PRIOR APPLICATION NUMBER: 60/091,183
; RIOR FILING DATE: 1998-06-30
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: PRELSEQ for Windows Version 4.0
; SEQ ID NO 32
; LENGTH: 15
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Pred. No. 3.8e+05;
1; Mismatches 0; Indels
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80.0%;
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Best Local Similarity Bu...
4; Conservative
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Best Local Similarity 50.0
Matches 5; Conservative
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APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Iwama, Akiko
Kino, Kohsuke
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6 QNKEDAWIEH 15
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10 DVWIDHNSLS 19
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                                           Gaps
                                                                                                                                                                                                                                                              APPLICANT: Rosenberg, Steven
APPLICANT: Bookenberg, Steven
APPLICANT: Bookenberg, Michael
TITLE OF INVENTION: Peptide Inhibitors of Urokinase Receptor
TITLE OF INVENTION: Activity
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron Corporation
STREET: 4560 Horton Street
CITY: Emeryville
STATE: CA
COUNTY: USA
ZIP: 94608
COMPUTER READABLE FORM:
MEDIUM TYPE: THOPPY disk
COMPUTER: IBM PC Compatible
COMPUTER: ISM PC COMPATA:
APPLICATION NUMBER: US/08/438,759
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 Score 30; DB 1; Length 17;
Pred. No. 2e+02;
2; Mismatches 3; Indels
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Pred. No. 2e+02;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/370,567
FILING DATE:
APPLICATION NUMBER: US/08/061,514
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Green, Grant D.
REGISTATION NUMBER: 31,259
REFERENCE/DOCKET NUMBER: 0941.001
TELECOMMUNICATION INFORMATION:
TELEFAX: 510-65-3542
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                             Sequence 5, Application US/08438759
Patent No. 5679782
   28.0%;
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LENGTH: 17 amino acids
TYPE: amino acid
TOPOLOGY: linear
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Best Local Similarity 50.0
Matches 5; Conservative
Query Match
Best Local Similarity 50.0
Matches 5; Conservative
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4 DLWMRHYPLS 13
                                                                             10 DVWIDHNSLS 19
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                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                 US-08-438-759-5
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RESULT 15
4.09-428-082B-198
5.8equence 198, Application US/09428082B
Fatent No. 6660843
GENERAL INFORMATION:
APPLICANT: FEIGE, ULRICH
APPLICANT: LIU, CHUAN-FA

```
APPLICANT: CHEETHAM, JANET C.

APPLICANT: BOONE, THOMAS CHARLES

FILE REFERENCE: A-527

CURRENT APPLICATION NUMBER: US/09/428,082B

CURRENT PELING DATE: 1999-10-23

CURRENT FILING DATE: 1999-10-23

NUMBER OF SEQ ID NOS: 1133

SOFTWARE: Patentin version 3.1

SEQ ID NO 198

LING PALICATION UKR ANTAGONIST PEPTIDE

CHENTY ARTHER SEATURE:

CHENTY MATCH

COURTENT FILING DATE: 1986-10-23

NUMBER OF SEQ ID NOS: 1133

SOFTWARE: PATORE:

COURTENT PRINCEMATION: UKR ANTAGONIST PEPTIDE

COURTENT INFORMATION: UKR ANTAGONIST PEPTIDE

COURTENT SORDE SIMILARITY SO.0%; Pred. No. 2e+02;

MATCHER S; CONSERVATIVE 2; Mismatches 3; Indels

ON 10 DVWIDHNSLS 19

DD 4 DLWMRHYPLS 13

Search completed: January 26, 2005, 16:08:25

JOB time: 15:6 secs
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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      296681
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1608061 seqs, 361289386 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Published_Applications_AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Listing first 45 summaries
                                                                                                                                                                                                OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 ASTGVTISNNHFFNHHKVML 20
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-202-464-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minimum DB seq length: 0 Maximum DB seq length: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                     Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		*			SUMMARIES	
Result No.	Score	Query Match	Query Match Length DB	DB	ΙD	Description
	74	68.5	15	14	US-10-354-240-52	Sequence 52, Appl
· ~	68	63.0	15	14	US-10-354-240-51	Sequence 51, Appl
ო	52	48.1	15	14	US-10-354-240-53	Sequence 53, Appl
4	45	41.7	15	14	US-10-354-240-50	Sequence 50, Appl
'n	39	36.1	20	14	US-10-005-530-24	Sequence 24, Appl
9	36	33.3	19	10	US-09-791-524-2	Sequence 2, Appli
1	36	33.3	19	17	US-10-821-544-17	17
œ	35	32.4	18	14	US-10-349-543-4	Sequence 4, Appli
σ	35	32.4	19	10	US-09-791-524-1	-1
10	34	31.5	0	15	US-10-107-532-1259	125
11	34	31.5	Q	15	US-10-107-532-1786	Sequence 1786, Ap
12	34	31.5	9	15	US-10-107-532-2854	Sequence 2854, Ap
13	34	31.5	6	15	US-10-107-532-4582	Sequence 4582, Ap

14	3.4	2,15	σ	4	118-10-107-532-4644	Semionre 4644 An
*	7	0.10	'n	7	7CC-/0T-0T-	-
12	34	31.5	10	12	-10-107-	1504,
16	34	31.5	10	15	US-10-107-532-2021	2021,
17	34	31.5	10	15	US-10-107-532-2613	2613,
18	34	31.5	20	15	7	3084,
19	34	31.5	70	15	-10-107-	3654,
20	34	31.5	10	15	US-10-107-532-5281	Sequence 5281, Ap
21	34	31.5	10	15	US-10-107-532-5294	5294,
22	34	31.5	10	15		5490,
23	34	31.5	15	15	US-10-107-532-5936	5936,
24	34	31.5	15	15	-10-107-53	5949,
25	34	31.5	15	15	∹	6009
26	34	31.5	15	12	-10-107-532	6009
27	34	31.5	15	15	-10-107-	6032,
28	34	31.5	15	12	US-10-107-532-6069	6909
29	34	31.5	15	12	-10-107-532	6095,
30	34	31.5	15	12	10-107-	6102,
31	34	31.5	15	15	-10-107-532	6113,
32	34	31.5	15	12	7	611
33	34	31.5	50	17	US-10-476-104-16	16,
34	32	29.6	σ	15	7	729,
35	32	29.6	10	14	031-	Sequence 88, Appl
36	32	29.6	10	15	US-10-107-532-394	394,
37	32	29.6	10	15	US-10-107-532-1481	148]
38	32	29.6	10	12	US-10-107-532-2074	207
39	32	29.6	10	12	US-10-107-532-5339	
40	32	29.6	10	15	US-10-107-532-5401	Sequence 5401, Ap
41	32	29.6	15	σ	US-09-729-835-116	116,
42	32	29.6	15	15	US-10-107-532-6013	
43	32	29.6	15	15	US-10-373-809-116	116
44	31	28.7	14	12	US-10-417-895A-33	33,
45	31	28.7	15	14	US-10-120-604-210	Sequence 210, App

## ALIGNMENTS

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GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Mame, Akinori
APPLICANT: Mame, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: SPO-103D1
ITILE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Diseareller Reference: SPO-103D1
CURRENT FILING DATE: 2003-01-29
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR PLING DATE: 1998-09-09
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 52
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 68.5%; Score 74; DB 14; Length 15; Best Local Similarity 93.3%; Pred. No. 0.00013; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (1). T(15); OTHER INFORMATION: Cryjl peptide, Figure 1, Row 38 US-10-354-240-52
Sequence 52, Application US/10354240 Publication No. US20030185847A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Cryptomeria japonica
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Gaps ö

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| APPLICANT: Sone, Toshio | APPLICANT: Kume, Akinori | APPLICANT: Kume, Akino | APPLICANT: Kino, Kohsuke | TITLE OF INVENTION: POPTIGE-Based Immunotherapeutic Agent for Treating Allergic Disea | FILE REFERENCE: SPO-103D1 | CURRENT APPLICATION NUMBER: US/10/354,240 | FRICR APPLICATION NUMBER: PCT/JP97/00740 | FRICR FILING DATE: 1997-03-10 | FRICR FILING DATE: 1998-09 | SPRICR FILING DATE: 1998-09 | SOGTWARE: Patentin version 3.1 | SEQ ID NO S: 174 | SEQ ID NO S: 1
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APPLICANT: Greenplate, John T.
APPLICANT: Greenplate, John T.
APPLICANT: Purcell, John F.
APPLICANT: Purcell, John F.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CONTROLLING INSECTS
FILE REFERENCE: 11899. OOZ2. DVUSGO. (MOBT:022--2)
CURRENT APPLICATION NUMBER: 09/10/05,530
CURRENT FILING DATE: 2001-10-26
PRIOR FILING DATE: 1998-04-21
PRIOR FILING DATE: 1998-04-21
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin Version 3.1
SEQ ID NO 24
                              Gaps
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                                 Indels
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CCATION: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 36
US-10-354-240-50
  100.0%; Pred. No. 0.32;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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; OTHER INFORMATION: Synthetic Peptide
US-10-005-530-24
                                                                                                                                                                                                                                                                        ; Sequence 50, Application US/10354240; Publication No. US20030185847A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24, Application US/10005530 Publication No. US20030026795A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41.7%;
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                              9; Conservative
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                                                                                 12 FFNHHKVML 20
                                                                                                                                    2 FFNHHKVML 10
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Best Local Similarity
  Best Local Similarity
Matches 9; Conserv
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                                                                                                                         TITLE OF INVENTION PEPTIDE Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
PRIOR PLING DATE: 10903-01-29
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR PILING DATE: 1997-03-10
PRIOR PILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
0
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: MISC_FEATURE
CCATTON: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 37
US-10-354-240-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 68; DB 14;
Pred. No. 0.0011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 53, Application US/10354240
Publication No. US20030185847A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.0%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ASTGVTISNNHFFNH 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 86.7-
Local 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
1 TISNNLFFNHHKVML
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-354-240-53
                                                                            RESULT 2
US-10-354-240-51
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LENGTH: 15
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TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes FILE REFERENCE: A3319A

CURRENT APPLICATION NUMBER: US/09/791,524

CURRENT FILING DATE: 2001-02-2

PRIOR APPLICATION NUMBER: 60/09828

PRIOR APPLICATION NUMBER: 1998-08-27

NUMBER OF SEQ ID NOS: 150

SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                         Haney, David N.
Varga, Janos
TITLE OF INVENTION: CYCLIC PEPTIDES THAT BIND TO
UROKINASE-TYPE PLASMINOGEN ACTIVATOR RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: LIVNAT, SHMUEL
REGISTRATION NUMBER: 33,949
REFERENCE/DOCKET NUMBER: ANG-001/DIV (80144-0007)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 955-8787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32.4%; Score 35; DB 14; Length 18; 46.2%; Pred. No. 1.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: RADER, FISHMAN & GRAUER
STREET: 1233 20TH STREET NW, SUITE 501
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/349,543
FILING DATE: 22-Jan-2003
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/285,783
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                                                                        US-10-349-543-4
Sequence 4, Application US/10349543
Publication No. US20030166514A1
GENERAL INFORMATION:
APPLICANT: Jones, Terence R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09791524
Publication No. US20030143209A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 18 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 20036
COMPUTER READABLE FORM:
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| :|| :| | | S GTAVSNKYFSNLH 17
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                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 9
US-09-791-524-1
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                                                                                                     RESULT 8
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Sequence 2, Application US/09791524

Sequence 2, Application US/09791524

Fublication No. US2030143209A1

GENERAL INFORMATION:

APPLICANT: Aventis Pharmaceuticals Products Inc.

TITLE OF INVENTION:

TITLE OF INVENTION:

TITLE REFERENCE:

CURRENT APPLICATION NUMBER: US/09/791,524

CURRENT FILING DATE: 2001-02-22

PRIOR APPLICATION NUMBER: 60/09828

PRIOR FILING DATE: 1998-08-27

NUMBER OF SEQ ID NOS: 150

SOFTWARE: Patentin version 3.0
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| Sequence 17, Application US/10821544
| Publication No. US20040265797A1
| GENERAL INFORMATION:
| APPLICANT: ROSENBERG, STEVE
| APPLICANT: CHAPMAN, HAROLD
| TITLE OF INVENTION: PEPTIDE LIGANDS OF THE UROKINASE RECEPTOR
| FILE REFERENCE: 014024-0284105/10/821,544
| CURRENT APPLICATION NUMBER: US/09/155,260C
| PRIOR PPLICATION NUMBER: PCT/US97/05199
| PRIOR PLILING DATE: 1998-09-23
| PRIOR PLILING DATE: 1998-09-23
| PRIOR PLILING DATE: 1997-03-28
| PRIOR PLILING DATE: 1996-03-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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                     Score 39; DB 14; Length 20;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
                                                                           Indels
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Pred. No. 1.2e+02;
2; Mismatches 5;
                                                                           2; Mismatches
                        36.1%;
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ilarity 46.2%;
Conservative
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Best Local Similarity 46.2
Matches 6; Conservative
                        36.1
Best Local Similarity 54.5
Matches 6; Conservative
                                                                                                                                                                 2 VVLEQNNFFNH 12
                                                                                                                           5 VTISNNHFFNH 15
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CORGANISM: Adenovirus
US-09-791-524-2
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Best Local Similarity
Matches 6; Conserv
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LENGTH: 19
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4 GVTISNNHFFNHH 16

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### APPLICANT: MOITEON, NOTES APPLICANT: Afar, NOTES APPLICANT: Afar, Daniel E.H.
### APPLICANT: Afar, Daniel E.H.
### APPLICANT: Afar, Daniel E.H.
### APPLICANT: Raiteano, Arthur
APPLICANT: Raiteano, Arthur
APPLICANT: Challita-Eid, pia M.
### APPLICANT: Challita-Eid, pia M.
### APPLICANT: Challita-Eid, Pia M.
### APPLICANT: Challita-Eid, Pia M.
### APPLICANTON: Number: 10540-00
### APPLICATION NUMBER: 60/283,112
### APPLICATION NUMBER: 60/284,630
### APPLICATION NUMBER: 60/286,630
### RIUNG DATE: 2001-04-25
### APPLICATION NUMBER: 60/286,630
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 50.0%; Pred. No. 1.5e+06;
Matches 4; Conservative 3; Mismatches 1; Indels
FILE REFERENCE: 51158-200064.00
CURRENT APPLICATION NUMBER: US/10/107,532
CURRENT FILING DATE: 2002-08-05
PRIOR PILING DATE: 2001-04-10
PRIOR FILING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: 60/286,630
PRIOR FILING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 6321
SOFTWARE: PRAESEQ for Windows Version 4.0
SEQ ID NO 1786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Morrison, Karen Jane Meyrick
Morrison, Robert Kendall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; sequence 4582, Application US/10107532; Publication No. US20040003418A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-107-532-2854, Application US/10107532; Sequence 2854, Application US/10107532; Publication No. US20040003418A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Agensys, Inc.
APPLICANT: Jakobovits, Aya
APPLICANT: Faris, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Agensys, Inc. APPLICANT: Jakobovits, Aya
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                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapien
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NRFYSHHR 9
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2 NRFYSHHR 9
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Best Local Similarity
Matches 4; Conserv
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APPLICANT: Faris, Mary
APPLICANT: Faris, Mary
APPLICANT: Morrison, Robert Kendall
APPLICANT: Morrison, Robert Kendall
APPLICANT: Morrison, Robert Kendall
APPLICANT: Morrison, Robert Kendall
APPLICANT: Afar, Daniel E.H.
APPLICANT: Afar, Daniel E.H.
APPLICANT: Ge, Wangmao
APPLICANT: Ge, Wangmao
APPLICANT: Challita-Eid, Pia M.
ITLE OF INVENTION: Entitled 158P3D2 Useful in Treatment and Detection of Cancer;
ITLE OF INVENTION: Entitled 158P3D2 Useful in Treatment and Detection of Cancer;
ITLE OF INVENTION: Entitled 158P3D2 Useful in Treatment and Detection of Cancer;
ITLE OF INVENTION: MOMBER: 60/283,112
PRIOR FILING DATE: 2001-04-10
PRIOR PLING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 6321
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 1259
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APPLICANT: Raitano, Arthur
APPLICANT: Challita-Eal, Pia M.
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 158P3D2 Useful in Treatment and Detection of Cancer
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APPLICANT: Agensys, Inc.
APPLICANT: Jakobovits, Aya
APPLICANT: Faris, Mary
APPLICANT: Morrison, Raren Jane Meyrick
APPLICANT: Morrison, Robert Kendall
APPLICANT: Hubert, Rene S.
APPLICANT: Afar, Daniel B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-107-532-1786; Application US/10107532; Sequence 1786; Application No. US20040003418A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1259, Application US/10107532
Publication No. US20040003418A1
GENERAL INFORMATION:
APPLICANT: Agensys, Inc.
APPLICANT: Jakobovits, Aya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 50.0°
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                                                                                                                                                                                                                                                                                 Conservative
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US-10-107-532-1259
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                          ; SEQ ID NO 1
; LENGTH: 19
; TYPE: PRT
; ORGANISM: Adenovirus
US-09-791-524-1
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Matches 6; Conserva
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APPLICANT:
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APPLICANT: Hobert, Rene S.
APPLICANT: Hobert, Rene S.
APPLICANT: Hobert, Rene S.
APPLICANT: Ge, Wangmao
APPLICANT: Ge, Wangmao
APPLICANT: Challita-Eid, Pia M.
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 15893D2 Useful in Treatment and Detection of Cancer FILE REPERENCE: 51158-200064.00
CURRENT APPLICATION NUMBER: US/10/107,532
CURRENT PILING DATE: 2002-08-05
PRIOR FILING DATE: 2001-04-10
PRIOR FILING DATE: 2001-04-10
PRIOR FILING DATE: 2001-04-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 6321
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1504
LENGTH: 10
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Job time : 56.5 secs
                                                                                                                                                                                                                                                        Morrison, Karen Jane Meyrick
Morrison, Robert Kendall
                                                                               Sequence 1504, Application US/10107532 Publication No. US20040003418A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31.5%;
50.0%;
                                                                                                                                                            APPLICANT: Agensys, Inc.
APPLICANT: Jakobovits, Aya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 31.5
Best Local Similarity 50.0
Matches 4; Conservative
                                                                                                                                                                                                                                   Faris, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Homo sapien
US-10-107-532-1504
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                      APPLICANT: Afer, Daniel B.H.
APPLICANT: Ge, Wangmao
APPLICANT: Raltano, Arthur
APPLICANT: Challita-Eid, Pia M.
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
FILE REFRENCE: 51158-200664.00
CURRENT APPLICATION NUMBER: US/10/107,532
CURRENT APPLICATION NUMBER: 60/283,112
FRIOR APPLICATION NUMBER: 60/286,630
FRIOR APPLICATION NUMBER: 60/286,630
FRIOR APPLICATION NUMBER: 60/286,630
FRIOR FILING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 6321
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Mary
APPLICANT: Mary
APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Morrison, Robert Kendall
APPLICANT: Hubert, Rene S.
APPLICANT: Afar, Daniel E.H.
APPLICANT: Afar, Daniel E.H.
APPLICANT: Raitano, Arthur
APPLICANT: Raitano, Arthur
APPLICANT: Challita-Eid, Pia M.
TITLE OF INVENTION: Undleic Acid and Corresponding Protein
TITLE OF INVENTION: Entiled 158P3D2 Useful in Treatment and Detection of Cancer;
CURRENT APPLICATION NUMBER: US/10/107,532
CURRENT APPLICATION NUMBER: 60/283,112
PRIOR FILING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: 60/286,630
PRIOR FILING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 6321
LENGTH: 9
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Morrison, Karen Jane Meyrick
Morrison, Robert Kendall
Hubert, Rene S.
Afar, Daniel E.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4644, Application US/10107532
Publication No. US20040003418A1
GENERAL INFORMATION:
APPLICANT: Agensys, Inc.
APPLICANT: Jakobovits, Aya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 50.0
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapien
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ORGANISM: Homo sapien
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1 NRFYSHHR 8
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Gaps ; 0

Score 34; DB 15; Length 10; Pred. No. 1.2e+02; 3; Mismatches 1; Indels

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24, Appli
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                                                                January 26, 2005, 15:54:02; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score.of the result being printed, and is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-09-142-524D-51
US-09-142-524D-53
US-08-467-023-43
US-08-467-023-43
US-08-467-023-43
US-08-747-915-4
US-08-142-590B-4
US-08-142-590B-24
US-08-142-590B-24
US-08-142-590B-24
US-08-142-590B-23
US-08-145-590B-23
US-08-436-772-26
US-08-436-773-18
                                                                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
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US-09-181-816-2
US-09-285-783-3
                                                                                                                                                                                                            478139 segs, 66318000 residues
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                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                              - protein search, using sw model
                                                                                                                                               1 ASTGVTISNNHFFNHHKVML 20
                                                                                                                                                                       BLOSUM62
Gapop 10.0 , Gapext 0.5
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                                                                                                                       US-09-202-464-21
108
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Match Length
          Copyright
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Perfect score:
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No.
                                                                       Run on:
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## ALIGNMENTS

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US-8-467-023-44

US-8-467-023-44

US-8-467-023-44

US-8-467-023-44

US-90308

PARTELL NOW 6090308

PAPLICANT: POILOCK, JOAnne;

APPLICANT: Bond, Julian F;

APPLICANT: Bond, Julian F;

APPLICANT: Bond, Julian F;

APPLICANT: Brauer, Andrew;

APPLICANT: Brauer, Andrew;

APPLICANT: Brauer, Andrew;

APPLICANT: Powers; Steven P.

ITILE OF INVENTION: Japanese Cedar Pollen

NUMBERSEE: LIMMULGGIC PHARMACCHICAL COPPORATION, INC.

STREET: G10 Lincoln St.

CITY: Waltham

COUNTRY: USA

COUNTRY: USA

COUNTRY: USA

COUNTRY: USA

COUNTRY: USA

CONFUTER READABLE FORM:

MEDIUM TYPE: PLOPDY disk

COMPUTER: ENGINER FORM:

MEDIUM TYPE: PLOPDY disk

COMPUTER: IBM FC compatible

OPERATING SYSTEM: PC-0005/M9-DOS

SOFWARE: Date FORM:

MEDIUM TYPE: PLOPDY disk

COMPUTER: Date FORM:

MEDIUM TYPE: PLOPDY disk

COMPUTER: Date FORM:

MEDIUM TYPE: PLOPDY disk

COMPUTER: Date FORM:

MEDIUM TYPE: PLOPDY disk

CLASSIFICATION NUMBER: US/08/467,023

FILING DATE: June 6, 1995

APPLICATION NUMBER: 035-025

FILING DATE: December 6, 1994

MANGE: Jane E. Remillard

MANGE: Jane E. Re
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TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: US/09/0040
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn version 3.1
SEQ ID NO 53
LENGTH: 15
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APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mai-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Brauer, Andrew;
APPLICANT: Brauer, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immulogic Pharmaceutical Corporation, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match

48.1%; Score 52; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 9; Conservative 0; Mismatches 0; Indels
                                                               Length 15;
                                                                                                          Indels
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610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: MISC_FEATURE
COCATION: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 39
US-09-142-524D-53
OTHER INFORMATION: Cryj1 peptide, Figure 1, Row 37
                                                             63.0%; Score 68; DB 4; I
86.7%; Pred. No. 0.00018;
iive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                   Sequence 53, Application US/09142524D
Patent No. 6719976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 45, Application US/08467023
; Patent No. 6090386
GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
Garman, Richard D;
Kuo, Mei-Chang;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                          1 ASTGVTISNNHFFNH 15
                                                                                                                                                                                   Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 FFNHHKVML 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 FFNHHKVML 10
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STATE: MA
                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02154
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US-09-142-524D-53
                         ÚS-09-142-524D-51
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US-08-467-023-45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Mama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE REPRESENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1988-09-09
PRIOR FILING DATE: 1997-00-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 5.2
LENGTH: 15
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Best Local Similarity 93.3%; Pred. No. 2.1e-05;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                  86.1%; Score 93; DB 3; Length 20; 94.7%; Pred. No. 2.9e-08; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (1).7(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 38
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                                                                                                                                                                                                                                                                                       Sequence 52, Application US/09142524D Patent No. 6719976
GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
                                                                                                                                        2 STGVTISNNHFFNHHKVML 20
                                                                                                                                                                                   STGVTISNNLFFNHHKVML 20
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ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 TISNNHFFNHHKVML 20
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                                                      Query Match
Best Local Similarity 94.7
Matches 18; Conservative
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NAME/KEY: MISC_FEATURE
LOCATION: (1). (15)
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                                                                                                                                                                                                                                                       RESULT 2
US-09-142-524D-52
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             US-08-467-023-44
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                                                                                                                               APPLICANT: Griffeth, Irwin J.;
APPLICANT: Golock, Joanne;
APPLICANT: Bollock, Joanne;
APPLICANT: Garman, Richard D;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Exlay, Mark A;
APPLICANT: Exlay, Mark A;
APPLICANT: Exlay, Mark A;
APPLICANT: Fowers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Appanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Isaac, Barbara G.
APPLICANT: Greenplate, John T.
APPLICANT: Romano, Charles P.
APPLICANT: Romano, Charles P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CONTROLLING
TITLE OF INVENTION: INSECTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                 ImmuLogic Pharmaceutical Corporation, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 44; DB 3;
Pred. No. 1.6;
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; Sequence 24, Application US/09063733A
; Patent No. 6372211
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
                                                                  Sequence 43, Application US/08467023; Patent No. 6090386; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0%; Pi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 227-5941 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 STGVTISNN 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 610 LU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-467-023-43
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GENERAL INFORMATION:

APPLICANT: Kume, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Induma, Akiko

APPLICANT: Induma, Akiko

APPLICANT: Kino, Kohsuke

TITLE OF INVERNION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

FILE REFERENCE: SPO-103

CURRENT APPLICATION NUMBER: US/09/142,524D

CURRENT PILING DATE: 1998-09-09

PRIOR FILING DATE: 1997-03-10

NUMBER OF SEQ ID NOS: 174

SEQ ID NO 50

LENGTH: 15

LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     48.1%; Score 52; DB 3; Length 20; 100.0%; Pred. No. 0.085; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41.7%; Score 45; DB 4; Length 15; 90.0%; Pred. No. 0.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PLING DATE: June 6, 1995
FILING DATE: June 6, 1995
FILING DATE: June 6, 1995
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REPERRINCE/POCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7400
TELEPRX: (617) 227-7400
TELEPRX: (617) 227-7400
TELEPRX: (617) 227-75941
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (1).7(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 36
US-09-142-524D-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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Sequence 50, Application US/09142524D
Patent No. 6719976
                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 90.0
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 ASTGVTISNN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: MISC_FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 FFNHHKVML 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-467-023-45
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MSG-08-142-590B-4
| Sequence 4, Application US/08142590B
| Sequence 4, Application US/08142590B
| Sequence 4, Application US/08142590B
| Patent No. 612076:
| Patent No. 612076:
| APPLICANT: HIBINO, Tashihiko; TAKAHASHI, Tadahito; HORII, Izumi; and TITLE OF INVENTION: UROKINASE PLASMINOGEN ACTIVATOR FRAGMENTS
| TITLE OF INVENTION: UROKINASE PLASMINOGEN ACTIVATOR FRAGMENTS
| TITLE OF INVENTION: 25
| CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: LAHIVE & COCKFIELD, LLP STRET: 28 State Street
| CITY: Boston Massachusetts
| CONNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 3; Length 18; 36;
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                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                 Score 35; DB 2;
Pred. No. 36;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32.4%; Score 35; DB 46.2%; Pred. No. 36; tive 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXt
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/142,590B
FILING DATE: 25-OCT-1993
RPICR APPLICATION NUMBER: US 08/042,318
FILING DATE: 02-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: MYETS, Paul L.
REGISTRATION NUMBER: 35,965
REFERENCE/DOCKET NUMBER: 35,965
REFERENCE/DOCKET NUMBER: MGP-009CP
TELECOMMUNICATION INFORMATION:
                           TELERAX: (202) 887-0763
TELEX: 90-4030 MRSNPOERSWSH
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 4:
             (202) 887-1500
                                                                                                                                                                                                                                                     32.4%;
illarity 46.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 GVTISNNHFFNHH 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTCVSNKYFSNIH 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
; TOPOLOGY: linear
US-08-747-915-4
                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 6; Conserv
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US-08-142-590B-24
             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-142-590B-4
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APPLICANT: Jones, Terence R.
APPLICANT: Jones, David N.
APPLICANT: Haney, David N.
APPLICANT: Yarga, Janos
TITLE OF INVENTION: CYCLIC PEPTIDES THAT BIND TO
TITLE OF INVENTION: UNCKINASE-TYPE PLASMINOGEN ACTIVATOR RECEPTOR
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 PENNSYLVANIA AVENUE, NW
CITY: WASHINGTON
STATE: DC
COUNTRY: USA
ZIP: 20006-1812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36.1%; Score 39; DB 3; Length 20; ilarity 54.5%; Pred. No. 9.6; Conservative 2; Mismatches 1 Tanal
NUMBER OF SEQUENCES: 58

CORRESPONDENCE ADDRESS:
ADDRESSE: Arnold White & Durkee
STREET: PO Box 4433
CITY: Houston
STATE: TX
COUNTRY: USA
ZID: 77210-4433
COMPUTER READABLE FORM:
MEDLUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
OPERATIOR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CLASSIFICATION NUMBER: US/09/063,733A
FILING DATE: 21-APR-1998
CLASSIFICATION NUMBER: 33,062
REPERENCE/DOCKET NUMBER: 33,062
REPERENCE/DOCKET NUMBER: 33,062
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amin acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPULER: L.C. DOS/MS-DOS OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 SOFTWARE: Patentin Bales #1.0, Version #1.30 APPLICATION NDATA: US/08/747,915 FILING DATE: 12-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 12-NOV-1996
CLASSIFICATION: 514
ATORNEY AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
RECISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,959
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/08747915
Patent No. S942492
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 VTISNNHFFNH 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 VVLEQNNFFNH 12
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Matches 6; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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GOETINCK,
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Patent No. 6120765
GENERAL INFORMATION:
APPLICANT: HIBINO, Tashihiko; TAKAHASHI, Tadahito; HORII, Izumi; and TITLE OF INVENTION: UROKINASE PLASMINOGEN ACTIVATOR FRAGMENTS
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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               FILING DATE: 05-Apr-1999
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INPORMATION:
NAME: LIVNAT, SHWUEL
REGISTRATION NUMBER: 33,949
REFERENCE/DOCKET NUMBER: ANG-001/DIV (80144-0007)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 955-8787
TELEPAX: (202) 955-8781
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                         Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 3; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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Pred. No. 36;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 41;
2; Mismatches
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SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:

PEDING DATE: 25-0CT-1993
FILING DATE: 25-0CT-1993
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/042,318
FILING DATE: 02-APR-1993
ATTORNEY/AGENT INFORMATION:

NAME: Wyers, Paul L.

REGISTRATION NUMBER: 35,965
FELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEFAK: (617) 227-7400
TELEFAK: (617) 227-5941
INFORMATION FOR SEG ID NO: 23:
                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              32.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 32.4%;
Best Local Similarity 46.2%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GTCVSNKYFSNIH 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 46.2
Matches 6; Conservative
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 28 St
CITY: Boston
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                                                                   GOETINCK,
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                      Patent No. 6120765
GENERAL INFORMATION:
APPLICANT: HIBINO, TAShihiko; TAKAHASHI, Tadahito; HORII, Izumi; and TITLE OF INVENTION:
UROKINASE PLASMINOGEN ACTIVATOR FRAGMENTS
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/09285783

Patent No. 6514710

GENERAL INFORMATION:
APPLICANT: Jones, Terence R.
Haney, David N.
Varga, Janos

TITLE OF INVENTION: UROKINASE-TYPE PLASMINOGEN ACTIVATOR RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 35; DB 3; Length 18; Pred. No. 36; 2; Mismatches 5; Indels
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COMPUTER READABLE FORM:
MEDIUM TYBE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFWRARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: RADER, FISHWAN & GRAUER
STREET: 1233 20TH STREET NW, SUITE 501
CITY: WASHINGTON
STATE: DC
COUNTRY: USA
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/142,590B
FILING DATE: 25-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/042,318
FILING DATE: U2-APR-193
ATTORNEY/AGENT INFORMATION:

NAME: Myers, Paul L.

REGISTRATION NUMBER: 35,965
REFERENCE/DOCKET NUMBER: MGP-009CP
TELECOMMUNICATION INFORMATION:

TELECHIONE: (617) 227-7400
TELECHIONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                     ADDRESSEE: LAHIVE & COCKFIELD, LLP STREET: 28 State Street CITY: Boston STATE: Massachusetts COUNTRY: USA
Sequence 24, Application US/08142590B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 32.4
Best Local Similarity 46.2
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                           021:09
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                                                                                                                                                                                                                     APPLICANT: O'Rand, Michael G.
APPLICANT: Widgren, Esther E.
APPLICANT: Widgren, Richard T.
APPLICANT: Lea, Isabel
TITLE OF INVENTION: Sperm Antigen Corresponding to a
TITLE OF INVENTION: Sperm Antigen Corresponding to a
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26, Application US/08436772
Patent No. 5814456
GENERAL INFORMATION:
GAPLICANT: Widgren, Esther E.
APPLICANT: Richardson, Richard T.
APPLICANT: Richardson, Richard T.
APPLICANT: Sperm Antigen Corresponding to a Sperm TITLE OF INVENTION: Sperm Antigen Corresponding to a Sperm VITLE OF INVENTION: Zona Binding Protein Autoantigenic Epitope NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenneth D. Sibley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
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Pred. No. 39;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER REABABLE FORM:
MEDIUW TYPE: Floppy disk
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPPEARING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/166,195A
FILING DATE: UDEC 1993
CLASSIFICATION: 435
ATTONENY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665
REGISTRATION NUMBER: 31,665
REGISTRATION NUMBER: 31,665
RELEPHONE: 919-881-3140
TELEPHONE: 919-881-3175
INFORMATION POR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
TENGTH: 10 amino acids
                                                                                                                                                      Sequence 26, Application US/08166195A Patent No. 5480799 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Kenneth D. Sibley STREET: P.O. Box 34009 CITY: Charlotte STATE: No. 5480799th Carolina COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30.6%;
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Best Local Similarity 55.6
Matches 5; Conservative
                          4 GTCVSNKYFSNIH 16
GVTISNNHFFNHH 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 NNHFFNHHK 17
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                                                                                        COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/436,772
FILNG DATE: OB-MA-1995
CLASSIFICATION: 435
ATTONEY/AGBAT INFORMATION:
NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665
REBERRICE/POCKET NUMBER: 5470-73B
TELEFHONE: 919-881-3140
TELEFAX: 919-881-3175
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Search completed: January 26, 2005, 16:08:25 Job time : 14.6 secs
STREET: P.O. CC. CITY: Charlotte STATE: No. 5814456th Carolina Carmer: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 NNHFFNHHK 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 NNHAFQEHE 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: si
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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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| Cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.ppp:*
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| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.ppp:*
| Cgn2_6/ptodata/1/pubpaa/US10D_PUBC
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1608061 segs, 361289386 residues
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. Maximum Match 100%
. Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Published Applications AA:*
                                                                                                                                                                                                       OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-202-464-22
115
1 HFFNHHKVMLLGHSDIYSDD 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BLOSUM62
Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                 Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

					SUMMARIES	
Result No.	Score	% Query Match	% Query Match Length DB	DB	a.	Description
	94	66.1	15	14	US-10-354-240-53	
7	72	62.6	15	14	US-10-354-240-54	Sequence 54, Appl
3	52	45.2	15	14	US-10-354-240-52	25
4	46	40.0	15	14	US-10-354-240-55	Sequence 55, Appl
S	33	28.7	18	σ	US-09-864-761-34373	Sequence 34373, A
9	32	27.8	6	σ	US-09-839-497A-11	Sequence 11, Appl
7	32	27.8	6	14	US-10-354-698-11	Sequence 11, Appl
80	32	27.8	12	17	US-10-775-965-75	Sequence 75, Appl
σ	32	27.8	18	σ	US-09-734-520-36	Sequence 36, Appl
10	32	27.8	18	13	US-10-012-034A-36	Sequence 36, Appl
11	32	27.8	20	15	US-10-420-564-6	Sequence 6, Appli
12	31	27.0	• •	10	US-09-880-748-2968	Sequence 2968, Ap
13	31	27.0	17	14	US-10-293-418-2968	Sequence 2968. Ap

Semionce 32 anni	1045,	6	1045,	1046,	1045,	1046,	1045,		Sequence 1045, Ap			_	5, Aj	129	19,	4	4	-	3,	ñ	736,	13,	13,	17	18	22		22	22	38	equence 39, Appl
-33									10	9	<sub>G</sub>	y	S	6				34		-735	-736					7	8	221	222	38	-39 Seq
5 TIG-10-414-583.	ns-	-10-	us-	us-	us-	us-	ns-	ns-	US-10-666-	US-1	US-10-653-		US-08-981-490B-5			6 US-10-775-640			US-08-775-765-3	4 US-10-226-629A	4 US-10-226-629A-7	US-09-839-497A-1	ns-		us-	ns-	5 US-10-468-370	US-1	US-10	US-1	US-10-033-662
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7.0		31 27																					6	6	6	6	29 25	6	6	6	6
14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	. 38	39	40	41	42	43	44	45

## ALIGNMENTS

```
US-10-34-240-35

Sequence 53, Application US/10354240

Publication No. US20030185847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Nume, Akinori

APPLICANT: Iwama, Akinori

APPLICANT: Iwama, Akinori

APPLICANT: Iwama, Akino

APPLICANT: Spo-103b1

FILE REFERENCE: SPO-103b1

CURRENT PILING DATE: 1999-03-02-29

FRIOR APPLICATION NUMBER: US/10/354,240

FRIOR APPLICATION NUMBER: US/09/142,524

FRIOR APPLICATION NUMBER: US 09/142,524

FRIOR FILING DATE: 1999-09-09

WINDHER OF SEQ ID NOS: 174

SEQ ID NOS: 174

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SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SERVINE: PREFERENCE: SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

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SEQ ID N
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APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akino
APPLICANT: Iwama, Akino
APPLICANT: Iwama, Akino
APPLICANT: Iwama, Akino
TITIE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITIE OF INVENTION PEPTIGE : 1891-0129
CURRENT APPLICATION NUMBER: US/10/354,240
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-9-864-761-54373, Application US/09864761

j Sequence 34373, Application US/09864761

j Patent No. US20020048763A1

j GENERAL INFORMATION:

APPLICANT: Penn, Sharron G.

APPLICANT: Hank, David R.

APPLICANT: Hank, David R.

APPLICANT: Hank GENE EXPRESSION ANALYSIS BY MICROARRAY

TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY

TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY

TITLE OF INVENTION: UNMBER: US/09/864,761

CURRENT APPLICATION NUMBER: US 60/180,312

PRIOR PILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR PILING DATE: 2000-08-26

PRIOR PILING DATE: 2000-08-03

PRIOR APPLICATION NUMBER: US 60/232,366

PRIOR PILING DATE: 2000-09-04

PRIOR PILING DATE: 2000-09-07

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2001-01-30
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                              Indels
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OTHER INFORMATION: Cryj1 peptide, Figure 1, Row 41
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; Pred. No. 0.61;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 46;
Pred. No. 4
                                                                                                                                                                                                                                                                     Sequence 55, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Cryptomeria japonica
FEATURE:
  Similarity 100.0%; 9; Conservative C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            40.0%;
80.0%;
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                                                                                   2 FFNHHKVML 10
                                                                                                                  7 FFNHHKVML 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: MISC FEATURE
  Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-354-240-55
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LENGTH: 15
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APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Lauma, Akinor
APPLICANT: Iwama, Akino
APPLICANT: Iwama, Akino
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103D1
CURRENT FILING DATE: 1093-01-29
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SEQ ID NO S: 174
SEQ ID NO S: 174
                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT FILING DATE: 2003-01-29
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO SEQ ID NOS: 174
LENGTH: 15
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86.7%; Pred. No. 0.00065;
tive 0; Mismatches 2; Indels
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COTHER INFORMATION: Cryjl peptide, Figure 1, Row 38
US-10-354-240-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (1). [(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 40
US-10-354-240-54
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                                                                                                                                     sequence 54, Application US/10354240
publication No. US20030185847A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Cryptomeria japonica
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           2 FFNHHKVMLLGHDD 15
                                                                                                                                                                                                                                               APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
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NAME/KEY: MISC FEATURE
                                                                                                                                                                                                                         APPLICANT: Sone, Toshio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: MISC_FEATURE
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Best Local Similarity
Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 3
US-10-354-240-52
                                                                                                                  US-10-354-240-54
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Sequence 11. Application US/10354698
| Publication No. US20030186416A1
| GENERAL INFORMATION:
| APPLICANT: Pallas, David C. |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase, |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase |
| TITLE OF INVENTION: Coding Sequence for Protein Phosphatase, |
| TITLE OF INVENTION NUMBER: 05/092,202 |
| PRIOR FILING DATE: 1999-04-16 |
| PRIOR FILING DATE: 1999-04-16 |
| PRIOR FILING DATE: 2001-04-16 |
| PRIOR FILING DATE: 2001-04-16 |
| PRIOR FILING DATE: 2001-04-16 |
| SEQ ID NOS: 17 |
| SOFTWARE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| TENTANTION |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
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| PRIOR FILING DATE: Patentin Ver. 2.0 |
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| PRIOR FILING DATE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
| SEQ ID NO 11 |
| PRIOR FILING DATE: Patentin Ver. 2.0 |
| PRIOR FILING DATE: P
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US-10-775-965-75

Sequence 75, Application US/10775965

Publication No. US20040209808A1

GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
APPLICANT: Rornacker, Michael
TITLE OF INVENTION: MODULATORS OF HUMAN G-PROTEIN COUPLED RECEPTORS
FILE REPERENCE: D0286 NP
CURRENT APPLICATION NUMBER: US/10/775,965
CURRENT FILING DATE: 2004-02-10
PRIOR FILING DATE: 2003-02-11
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Pred. No. 1.5e+06;
2; Mismatches 0; Indels
                                                                                                                                                               Length 9;
                                                                                                                                                           27.8%; Score 32; DB 9; I
71.4%; Pred. No. 1.5e+06;
7ative 2; Mismatches 0;
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Best Local Similarity 71.4%;
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SOFTWARE: Patentin version 3.2
SEQ ID NO 75
LENGTH: 12
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                                                                                                                                                           Query Match 27.8
Best Local Similarity 71.4
Matches 5; Conservative
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                                                        ; ORGANISM: Homo sapiens
US-09-839-497A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                 8 VMLLGHS 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8 VMLLGHS 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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1 IMLIGHS 7
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                        TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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N: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5

N: EXPRESSED IN LUNG, SIGNAL = 1.3

N: EXPRESSED IN HELA, SIGNAL = 1.1

N: EXPRESSED IN PLACENTA, SIGNAL = 1.5

N: EXPRESSED IN PLACENTA, SIGNAL = 1.3

N: EXPRESSED IN BONE MARROW, SIGNAL = 1.3

N: EXPRESSED IN HEART, SIGNAL = 1.3

N: EXPRESSED IN HEART, SIGNAL = 1.3
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PRIOR APPLICATION NUMBER: PCT/USO1/00667

PRIOR FILING DATE: 2001-01-30

PRIOR PLILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PLILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00669

PRIOR APPLICATION NUMBER: PCT/USO1/00665

PRIOR APPLICATION NUMBER: PCT/USO1/00668

PRIOR APPLICATION NUMBER: PCT/USO1/00668

PRIOR PLILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00661

PRIOR PLILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: US 09/608,408

PRIOR FILING DATE: 2000-09-21

PRIOR FILING DATE: 2000-06-30

PRIOR FILING DATE: 2000-06-30

PRIOR PLILING DATE: 2000-06-30

PRIOR PLILING DATE: 2000-06-30

PRIOR PLILING DATE: 2001-01-29

NUMBER OF SEQ ID NOS: 49117

SEQ ID NOS: 49373

LENGTH: 18
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Pred. No. 4.9e+02;
4; Mismatches 4;
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OTHER INFORMATION: EXPRESSED IN
US-09-864-761-34373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 28.7%;
Similarity 33.3%;
4; Conservative
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7 HHHHHHVITIG 18
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Best Local Similarity
Matches 4; Conserv;
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Sequence 6, Application US/10420564

Sequence 6, Application US/10420564

Publication No. US20040001819A1

GENERAL INFORMATION:

APPLICANT: Bolen, Paul L.

APPLICANT: Cihak, Paul L.

APPLICANT: Cihak, Paul L.

TITLE OF INVENTION: Recombinant Kid Pregastric Esterase and Methods for TITLE OF INVENTION: Recombinant Kid Pregastric Esterase and Methods for TITLE OF INVENTION: Broadction and Use FILE REFERENCE: IFF-0009

TITLE OF INVENTION: WINBER: US/10/420,564

CURRENT APPLICATION NUMBER: US/10/043,665B

PRIOR FILING DATE: 2002-09-24

PRIOR FILING DATE: 1998-11-05

NUMBER OF SEQ ID NOS: 7

SOFTWARE: PatentIn Ver. 2.0

LENGTH: 20
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iSequence 2968, Application US/09880748

iSequence 2968, Application US/09880748

iSequence 2968, Application US/09880748

iPublication No. US2003005993741

iGENERAL INFORMATION:

APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REFREENCE: PP523

CURRENT APPLICATION NUMBER: US/09/880,748

CURRENT FILING DATE: 2000-06-15

PRIOR APPLICATION NUMBER: 60/210,210

PRIOR FILING DATE: 2000-10-17

PRIOR APPLICATION NUMBER: 60/276,248

PRIOR FILING DATE: 2001-03-16

PRIOR PRILING DATE: 2001-03-21

PRIOR APPLICATION NUMBER: 60/293,499

PRIOR PILING DATE: 2001-05-25

NUMBER OF SEQ ID NOS: 3239

SOFTWARE: PatentIN Ver. 2.0

INNORTH: 17
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27.8%; Score 32; DB 15; Length 20
Best Local Similarity 40.0%; Pred. No. 7.7e+02;
Matches 8; Conservative 1; Mismatches 9; Indels
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ORGANISM: Artificial Sequence
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Best Local Similarity 50.0
Matches 6; Conservative
7 FICHHVVRLLG 17
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US-09-880-748-2968
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Publication No. US20020137141A1

Publication No. US20020137141A1

Publication No. US20020137141A1

APPLICANT: Ben-Sasson, Shmuel

TITLE OF INVENTION: SHORT PEPTIDES FROM THE A-REGION OF

TITLE OF INVENTION: ACTIVITY

FILE REPERENCE: BEN-SASSON=5A

CURRENT APPLICATION NUMBER: US/10/012,034A

CURRENT APPLICATION NUMBER: US/10/012,034A

CURRENT PILING DATE: 2001-12-11

PRIOR FILING DATE: 2000-12-11
                                                                                                                                                                                            Sequence 36, Application US/09734520

Batent No. US20020115173A1

BATELICANT: Ben-Sasson, Shmuel
TITLE OF INVENTION: SHORT PEPTIDES FROM THE A-REGION OF
TITLE OF INVENTION: ACTIVITY
FILLE REFERENCE: 1242.2003-000
CURRENT APPLICANTION NUMBER: US/09/734,520
CURRENT PELING DATE: 2000-12-11
NUMBER OF SEQ ID NOS: 122
SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 36
LENGTH: 18
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      Indels
         3;
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Pred. No. 6.9e+02;
0; Mismatches 4.
         Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 36
LENGTH: 18
TYPE: PRT
ORGANISM: Artificial Sequence
         2;
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Best Local Similarity 63.6
Matches 7; Conservative
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Best Local Similarity 63.6
Matches 7; Conservative
           5; Conservative
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                                                     4 NHHKVMLLGH 13
                                                                               3 NSHKIWMLPH 12
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; LOCATION: (0)...(18)
US-10-012-034A-36
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                                                                                                                                                                RESULT 9
US-09-734-520-36
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4; Indels

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                                                                                                                                                                                                                                                                                                                                  APPLICANT: EIGE, ULRICH
APPLICANT: EIGE, ULRICH
APPLICANT: LIU, CHUAN FA
APPLICANT: DOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
FILE REPERENCE: A-57
CURRENT APPLICATION NUMBER: US/10/609,217
CURRENT FILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: US/09/428,082B
PRIOR FILING DATE: 1999-10-22
PRIOR APPLICATION NUMBER: 60/105,371
PRIOR PLING DATE: 1999-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE: PATENTI Version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20
  Indels
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Pred. No. 1.1e+03;
Mismatches
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                                                                                                                                                                                                                                                  Sequence 1045, Application US/10609217
Publication No. US20040044188A1
GENERAL INFORMATION:
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                                                        4 NHHKV----MLLGHSD 15
                                                                                                           2 SHOKVTLLOLLLGHKN 17
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ORGANISM: Artificial Sequence
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7; Conservative
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7 FLPHYKNTSLGH 18
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Best Local Similarity
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Matches
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Sequence 33, Application US/10414583

Publication No. US20040005636A1

GENERAL INFORMATION:

APPLICANT: Guy, Mondey Kiplin

APPLICANT: Geistlinger, Timothy Ross

TITLE OF INVENTION: TO A PROTEIN

FILE REFERENCE: 9811-015-999

CURRENT APPLICATION NUMBER: US/10/414,583

CURRENT FILING DATE: 2003-04-15

PRIOR FILING DATE: 2002-04-15
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                                                                                                                                                                                                           Sequence 2968, Application US/10293418
Publication No. US2003022395641
GENERAL INFORMATION:

APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REPRENCE: PS23P2
CURRENT APPLICATION NUMBER: US/10/293,418
FRIOR APPLICATION NUMBER: 00/331,469
PRIOR FILING DATE: 2002-11-16
PRIOR PPLING DATE: 2001-12-19
PRIOR PLILING DATE: 2001-12-19
PRIOR PLILING DATE: 2001-06-15
PRIOR PLILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR PLILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR PLILING DATE: 2001-03-16
PRIOR PLILING DATE: 2001-03-16
PRIOR PPLING DATE: 2001-03-16
PRIOR PPLICATION NUMBER: 60/210,210
PRIOR PPLING DATE: 2000-01-17
PRIOR PLING DATE: 2000-01-17
PRIOR PPLING DATE: 2000-01-17
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Pred. No. 9.1e+02;
2; Mismatches 4; Indels
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SOFTWARE: PatentIn version 3.2
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Best Local Similarity 50.0
Matches 6; Conservative
                       20
                                                  9 MLLGHSDIYSDD 20
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5 ILTGYSDYYGMD 16
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US-10-293-418-2968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Homo sapiens
US-10-414-583-33
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Best Local Similarity
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SEQ ID NO 33
LENGTH: 20
TYPE: PRT
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ALIGNMENTS

Appl Appl Appl Appl Appl Appl

Sequence sequence Sequence Sequence Sequence Sequence Sequence Sequence ( Sequence Sequence (

Sequence 1 Sequence 4 Sequence 1

Sequence Sequence

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Patent No. 600386
GENERAL INFORMATION:
APPLICANT: Polick, Joane;
APPLICANT: Bond, Julian F;
APPLICANT: Bond, Julian F;
APPLICANT: Bond, Julian F;
APPLICANT: Kuo, Mei-Chang D;
APPLICANT: Kuo, Mei-Chang B;
APPLICANT: Yeung, Siu-mei H;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Brauer, Andrew;
APPLICANT: Alergenic Proteins And Peptides From TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Spanese Cedar Pollen
NUMBER OF SEQUENCES: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: ImmuLogic Pharmaceutical Corporation, STREET: 610 Lincoln St CITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
US-09-019-346A-12

US-09-423-353-19

US-09-127-946-4

US-09-293-322C-14

US-09-89-497A-13

US-07-616-910-46

PCT-US91-08497-46

US-08-482-778-15

US-09-139-780A-103

US-09-440-781-76

US-09-440-781-76

US-09-440-781-76

US-08-408-604A-75

US-08-955-777-9

US-08-955-777-9

US-08-955-777-9

US-08-95-777-9

US-08-95-777-9

US-08-96-910-1

US-07-616-910-1

US-07-616-910-1
                                                                                                                                                                                                                                                                                                                           ; Sequence 45, Application US/08467023 ; Patent No. 6090386
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COUNTRY: USA
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US-08-467-023-45
    11, Appl
11, Appl
10, Appl
6, Appli
6, Appli
60, Appli
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1046, Ap
2, Appli
5, Appli
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                                                                                    January 26, 2005, 15:54:02; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
              GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
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US-08-542-051-21
US-09-428-082B-1045
US-09-458-082B-1046
US-08-945-848-2
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US-09-142-524D-54

US-09-142-524D-52

US-08-467-023-44

US-09-142-524D-55

US-09-121-211-4

US-09-233-32C-11

US-09-233-32C-11

US-09-121-211-10

US-09-121-211-10

US-09-121-211-10

US-09-121-211-10

US-09-121-211-10

US-09-131-211-10

US-09-131-211-10

US-09-131-211-10

US-09-131-211-10

US-09-131-211-10

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-08-530-524A-19
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Maximum Match 100%
Listing first 45 summaries
                                                               OM protein - protein search, using sw model
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115
1 HFFNHHKVMLLGHSDIYSDD 20
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Gapop 10.0 , Gapext 0.5
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Match Length
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Perfect score:
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APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea-
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea-
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disease.
TITLE OF INVENTION: 1994-09-09
CURRENT APPLICATION NUMBER: DCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn Version 3.1
SEQ ID NO 52
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APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Appanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:

| FEATURE:
| NAME/KEY: MISC_FEATURE
| LOCATION: (1)...(15)
| OTHER INFORMATION: Cryjl peptide, Figure 1, Row 38
US-09-142-524D-52
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 40 US-09-142-524D-54
                                                                Score 72; DB 4;
Pred. No. 0.00015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45.2%; Score 52; DB 4;
100.0%; Pred. No. 0.14;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                     US-09-142-524D-52; Sequence 52, Application US/09142524D; Patent No. 6719976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 44, Application US/08467023; Patent No. 6090386; GENERAL INFORMATION: APPLICANT: Griffeth, Irwin J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Garman, Richard D, APPLICANT: Kuo, Mei-Chang; APPLICANT: Yeung, Siu-mei H.; APPLICANT: Brauer, Andrew; APPLICANT: Exley, Mark A.; APPLICANT: POWETE, Steven P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Cryptomeria japonica
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Bond, Julian F.;
                                                                   Query Match 62.6%;
Best Local Similarity 86.7%;
Matches 13; Conservative
                                                                                                                                                          6 HKVMLLGHSDIYSDD 20
                                                                                                                                                                                                1 HKVMLLGHDDAYSDD 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairrki, Kazuo
APPLICANT: Inama, Akiko
APPLICANT: Kino, Kohsuke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 FFNHHKVML 10
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COUNTRY:
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APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Wine, Akinori
APPLICANT: Inwama, Akinori
APPLICANT: Inwama, Akino
APPLICANT: Inwama, Akino
APPLICANT: Kino, Kohsuke
APPLICANT: Kino, Kohsuke
APPLICANT: Kino, Kohsuke
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: PCT/UP97/00740
PRIOR PILING DATE: 1997-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO S: 15
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Pred. No. 3.7e-05;
0; Mismatches 1; Indels
                                                 Query Match
Best Local Similarity 89.5%; Pred. No. 2.5e-08;
Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (1)...(15) ...
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 39 US-09-142-524D-53
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; Sequence 54, Application US/09142524D
; Patent No. 6719976
                                                                                                                                                                                                                                                                              Sequence 53, Application US/09142524D
Patent No. 6719976
                                                                                                                                          2 FFNHHKVMLLGHSDIYSDD 20
                                                                                                                                                                     2 FFNHHKVMLLGHDDAYSDD 20
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SOFTWARE: Patentin version 3.1
SEQ ID NO 54
IENGTH: 15
TYPE: PRT
ORGANISM: Cryptomeria japonica
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 FFNHHKVMLLGHSD 15
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                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairrki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: NAME/KEY: MISC_FEATURE LOCATION: (1). (15)
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                                                                                                                                                                                                                                                                     -09-142-524D-53
             US-08-467-023-45
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Gaps
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                                                                                                                                                                                APPLICANT: Giffeth, Irwin J.;
APPLICANT: Bollock, Joanne,
APPLICANT: Bollock, Joanne,
APPLICANT: Bond, Julian F.;
APPLICANT: Bornd, Richard D.;
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Tribe OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Allergenic Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 3; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ImmuLogic Pharmaceutical Corporation, Inc.
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APPLICANT: Shinohara, Toshimichi
APPLICANT: Shinohara, Direndra P.
APPLICANT: Chylack, Leo T.
TITLE OF INVENTION: Lens Epithelial Cell Derived Growth
TITLE OF INVENTION: Pactor
FILE REFERENCE: B0801/7116
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SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995

CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994

ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              025.6 USD2 (IMI-028CPD2)
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40.0%; Score 46; DB:
Best Local Similarity 80.0%; Pred. No. 1.5;
Matches 8; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                   ; Sequence 46, Application US/08467023
; Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-121-211-4
; Sequence 4, Application US/09121211
; Patent No. 6750052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET UNBER: 02
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 227-5941 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TREET: 610 Lincoln St. ITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 LGHSDIYSDD 20
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APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Twame, Akinori
APPLICANT: Iwama, Akinori
APPLICANT: Iwama, Akino
FILE REPREBUCE: SPO-103
FILE REPREBUCE: SPO-103
FURENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: PCT/JP97/00740
FRIOR APPLICATION NUMBER: PCT/JP97/00740
FRIOR APPLICATION NUMBER: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn version 3.1
SEQ ID NO SECOND 
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                                          MEDIUM TYPE: FLOPPY GIBK
COMPUTER: PLOPPY GIBK
COMPUTER: THOM COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTEN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: Unne 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGETY INFORMATION:
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEBRANCE: (617) 227-7400
TELEBRANCE: (617) 227-7400
TELEBRANCE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:

; LOCATION: (1).. (15)
; OTHER INFORMATION: Cryj1 peptide, Figure 1, Row 41
VS-09-142-224b-55
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Best Local Similarity 80.0
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
COMPUTER READABLE FORM:
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Best Local Similarity
Matches 9; Conserva
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TYPE: PRT
CORGANISM: Homo sapiens
US-09-839-497A-11
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US-09-121-211-10
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1 IMLIGHS 7
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NAME/KEY: PEPTIDE
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US-09-186-489-6
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LENGTH: 9
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## Sequence 11, Application US/09839497A
## Patent No. 6528295
## Septembral INFORMATION:
## APPLICANT: Pallas, David C.
## APPLICANT: Du, Xianxing
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Nathylesterase,
## TITLE OF INVENTION: Coding Sequence for Protein Phosphatase
## TITLE OF INVENTION: UNWERS: 60/082,202
## PRIOR FILING DATE: 2001-04-20
## PRIOR FILING DATE: 1999-04-16
## PRIOR FILING DATE: 1999-04-16
## SOFTWARE: PatentIN Ver. 2.0
## SOFTWARE: PatentIN Ver. 2.0
## SEQ ID NO 11
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US-09-293-322C-11
Sequence 11, Application US/09293322C
Facquence 11, Application US/0929332C
Fatent No. 623210
GENERAL INFORMATION:
APPLICANT: Pallas, David C
TITLE OF INVENTION: Coding Sequence for Protein Phosphatase Methylesterase,
TITLE OF INVENTION: Recombinant DNA Molecules and Methods
TITLE OF INVENTION Recombinant DNA Molecules and Methods
FILE REPREBUGE: 105-97
CURRENT APPLICATION NUMBER: US/09/293,322C
CURRENT PRILING DATE: 1999-04-16
PRIOR FILING DATE: 1998-04-16
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CURRENT APPLICATION NUMBER: US/09/121,211
CURRENT FILING DATE: 1998-07-23
EARLIER APPLICATION NUMBER: U.S. 60/053,549
EARLIER FILING DATE: 1997-07-23
NUMBER OF SEQ ID NOS: 22
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 4
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                               2 FFNHHKVMLLGHSDIY 17
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                                                                                                                                                                                                                                                                                                    33.94
Best Local Similarity 43.84
Matches 7; Conservative
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Best Local Similarity 71.4
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-293-322C-11
                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo Sapiens
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SOFTWARE: Patentin Ve
SEQ ID NO 11
LENGTH: 9
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1 IMLIGHS 7
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US-09-839-497A-11
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Sequence 6, Application US/09186489;
Patent No. 6375947;
GENERAL INFORMATION:
APPLICANT: Bolen, Paul L.
APPLICANT: GIARK, Paul L.
APPLICANT: Scharpf Jr., Lewis G.
TITLE OF INVENTION: Purified Recombinant Kid Pregastric Esterase, and
TITLE OF INVENTION: Processes for its Production and Use;
FILE REFERENCE: 5499/3 unwhere; US/09/186,489;
CURRENT FILING DATE: 1998-11-05;
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shingh, Dhirendra P.
APPLICANT: Shingh, Dhirendra P.
APPLICANT: Shingh, Dhirendra P.
APPLICANT: Chylack, Leo T.
TITLE OF INVENTION: Lens Epithelial Cell Derived Growth;
TITLE OF INVENTION: Pactor
FILE REFERENCE: B0801/7116
CURRENT FILING DATE: 1998-07-23
EARLIER APPLICATION NUMBER: U.S. 60/053,549
EARLIER PILING DATE: 1997-07-23
NUMBER OF EXQ ID NOS: 22
SOFTWARE: FastSEQ for Windows Version 3.0
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 32; DB 4; Length 15;
Pred. No. 1.4e+02;
1; Mismatches 7; Indels
Score 32; DB 4; Length 9;
Pred. No. 3.8e+05;
2; Mismatches 0; Indels
                                                     0; Indels
                                                                                                                                                                                                                                                                US-09-121-211-10
; Sequence 10, Application US/09121211
Patent No. 6750052
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 42.9%;
Matches 6; Conservative
     Query Match 27.8%;
Best Local Similarity 71.4%;
Matches 5; Conservative
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Gaps

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Patent. No. 5854387
GENERAL INFORMATION:
APPLICANT: Urry, Dan W.
APPLICANT: Xu, Jie
TITLE OF INVENTION: A Simple Method for the Purification of TIMER OF INVENTION: a Bioelastic Polymer
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB 1; Length 20;
Pred. No. 2.7e+02;
1; Mismatches 2; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/543,020
FILING DATE: 13-0CT-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Hughes, Melya J.
REGISTRATION NUMBER: 38,696
REGISTRATION NUMBER: BEL-016/01US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415 853 5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB 2; Ler
Pred. No. 2.7e+02;
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-293-1404
TELEFAX: 202-872-0493
TELEX: 440 069 AIS UI
INFORMATION FOR SEQ ID NO: 60: SEQUENCE CHARACTERISTICS:
LENGTH: 20 aming acids
                                                                                                                                                                                                                                                                                                                                                                                   27.0%;
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33.3%;
         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                Query Match 27.0
Best Local Similarity 66.7
Matches 6; Conservative
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ZIP: 94306-2155
COMPUTER REDABLE FORM:
MEDIUM TYPE: Florm:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
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US-07-678-974D-60
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                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/10043665B
; Sequence 6, Application US/10043665B
; GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Recombinant Kid Pregastric Esterase and Methods for Its
TITLE OF INVENTION: Production and Use
TITLE OF INVENTION: Production and Use
TITLE OF INVENTION: Production and Use
CURRENT FILING DATE: 1900-24
FRIOR APPLICATION NUMBER: US 09/186, 489
FRIOR FILING DATE: 1998-11-05
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 20
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Patent No. 5629146
GENERAL INFORMATION:
APPLICANT: DILLNER, JOAKIM
APPLICANT: DILLNER, LENA
TITLE OF INVENTION: METHOD FOR DETECTION OF HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: A modified OTHER INFORMATION: polyHis-enterokinase polypeptide sequence
                                                                    5,
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27.8%; Score 32; DB 3; Length 20; ilarity 40.0%; Pred. No. 1.9e+02; Conservative 1; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 27.8%; Score 32; DB 4; Length 20; Best Local Similarity 40.0%; Pred. No. 1.9e+02; Matches 8; Conservative 1; Mismatches 9; Indels
                                                             Indels
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CORRESPONDENCE ADDRESS:
ADDRESSE: 1720 RHODE ISLAND AVENUE, N.W.,
STREET: 170 RHODE ISLAND AVENUE, N.W.,
STATE: 0.C.
COUNTRY: U.S.A.
ZIP: 20036-3186
COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/07/678,974D
FILING DATE: 25-JUN-1991
                                                                                                                        1 HFFNHHKVMLLGHSDIYSDD 20
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ALISENBERG, Irin M.
REGISTRATION NUMBER: 19,007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                             Best Local Similarity
Matches 8; Conserv
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   Query Match
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Gaps

Search completed: January 26, 2005, 16:08:26 Job time : 15.6 secs

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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

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17: /cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1608061 segs, 361289386 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                        OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                        1 LGHSDIYSDDKSMKVTVAFN 20
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Gapop 10.0 , Gapext 0.5
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103
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                              Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

					SUMMARIES	
Result		* Query				
No.	Score	Match	Match Length DB	DB	D	Description
-	7.1	68.9	15	14	US-10-354-240-56	Sequence 56, Appl
8	69	67.0	15	14	US-10-354-240-55	Sequence 55, Appl
m	48	46.6	15	14	US-10-354-240-9	
4	48	46.6	15	14	US-10-354-240-57	Sequence 57, Appl
ß	48	46.6	15	14	US-10-354-240-158	158,
9	46	44.7	15	14	US-10-354-240-54	Sequence 54, Appl
7	32.5	31.6	20	10	US-09-853-079-218	
æ	32.5	31.6	20	15	US-10-294-443-218	218,
0	32	31.1	18	14	US-10-084-813-172	172,
10	32	31.1	20	14	US-10-212-679-399	399,
11	32	31.1	20	14	US-10-212-679-400	400,
12	32	31.1	20	15	US-10-079-137B-399	
13	32	31.1	20	15	US-10-079-137B-400	

Sequence 38, Appl Sequence 39, Appl Sequence 233, App Sequence 4, Appli Sequence 4, Appli Sequence 65, Appli	00440	120 4 4 6	40 -HWOH	5, Ap 38, A 169, 185, 401, 401, 19, A
14 US-10-033-741-38 14 US-10-033-662-39 14 US-10-285-394-233 16 US-10-775-640-4 16 US-10-775-423-4 14 US-10-242-056-65	14 US-10-262-794A-65 17 US-10-776-013-657 15 US-10-380-147-45 15 US-10-436-715-422 10 US-09-832-464-22	9 US-09-966-955A-23 14 US-10-092-367-5 14 US-10-092-367-137 14 US-10-092-367-153 14 US-10-210-023-98	15 US-10-351-334-342 17 US-10-867-460-98 14 US-10-186-681-3 14 US-10-105-232-118 14 US-10-105-332-131 14 US-10-189-437-105 14 US-10-189-437-118	9 US-09-966-95A-35 14 US-10-092-367-169 14 US-10-092-367-185 14 US-10-12-679-401 15 US-10-079-137B-401 14 US-10-284-400-19 14 US-10-284-083-13 14 US-10-268-336-8
4 4 4 8 8 6	13	88888	18 10 10 10 10 10	411 200 80 80 80 80
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## ALIGNMENTS

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JUNEAU PAPLICATION NO. US/10354240

| Publication No. US/20030185847A1
| GENERAL INFORMATION:
| APPLICANT: Kume, Akinori
| APPLICANT: Wame, Akinori
| PRIOR PAPLICATION NUMBER: US/10/354,240
| PRIOR PAPLICATION NUMBER: US 09/142,524
| PRIOR APPLICATION NUMBER: US 09/142,524
| PRIOR APPLICATION NUMBER: US 09/142,524
| WUMBER OF SEQ ID NOS: 174
| SOFTWARE: Patentin Version 3.1
| SEQ ID NO 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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    i LOCATION: (1) .. (15)
    i OTHER INFORMATION: Cryjl peptide, Figure 1, Row 42
    MS-10-354-240-56

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: MISC_FEATURE LOCATION: (1)...(15)
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7 YSDDKSMKVTVAFN 20

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APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Wume, Akinori
APPLICANT: Wume, Akinori
APPLICANT: Iwama, Akiko
APPLICANTION: POPLIde-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR PILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 57
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Mana, Akino
APPLICANT: Iwama, Akiko
"
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APPLICANT: Iwama, Akiko
"
"
"
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disearence: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT PILING DATE: 2003-01-29
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn version 3.1
SEQ ID NO 158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FRATURE:
NAME/KEY: MISC FEATURE
LOCATION: (1)...(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 43
US-10-354-240-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/10354240
o. US20030185847A1
                                                                                                                                                 Sequence 57, Application US/10354240 Publication No. US20030185847A1
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COTHER INFORMATION: Figure 7, Row US-10-354-240-158
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ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Cryptomeria japonica
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Best Local Similarity 100.
Matches 10; Conservative
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1 KSMKVTVAFN 10
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                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Sone, Toshio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 158, Applion Publication No. US2, GENERAL INFORMATION
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Sequence 9, Application US/10354240

Publication No. US20030185847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Wame, Akinori

CURRENT FAILON: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

FILE REFERENCE: SPO-103D1

CURRENT FILING DATE: 2003-01-29

PRIOR APPLICATION NUMBER: US/10/354,240

CURRENT FILING DATE: 1997-03-10

PRIOR FILING DATE: 1998-09-09

PRIOR FILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174

SED ID NOS: 174
                                                                                                                                                                                                 APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Wime, Akinori
APPLICANT: Wime, Akinori
APPLICANT: Wime, Akino
APPLICANT: Wime, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION NUMBER: US/10/354,240
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 55
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: MISC_FEATURE
CCATTON: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 41
US-10-354-240-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 69; DB 14;
Pred, No. 0.00025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                               Sequence 55, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Cryptomeria japonica
US-10-354-240-9
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86.7%;
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       2 YSDDKSMKVTVAFN 15
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Best Local Similarity 100.0
Matches 10; Conservative
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Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                         US-10-354-240-55
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US-10-354-240-9
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; Sequence 172, Application US/10084813
; Publication No. US20030068615A1
; Publication No. US20030068615A1
; GENERAL INFORMATION:
; APPLICANT: SAXINGER, CARL
; TITLE OF INVENTION: POLYPEPTIDES THAT BIND HIV GP120 AND RELATED NUCLEIC
; TITLE OF INVENTION: ACIDS, ANTIBODIES, COMPOSITIONS, AND METHODS OF USE
; FILE REPRENCE: 215875
; CURRENT APPLICATION NUMBER: US/10/084,813
; CURRENT FILING DATE: 2002-02-27
; PRIOR FILING DATE: 2000-08-25
; PRIOR FILING DATE: 1999-08-27
; NUMBER OF SEQ ID NOS: 1242
; SOFTWARE: Patentin Version 3.1
; SEQ ID NO 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: binding peptide US-10-084-813-172
                                                   Gaps
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                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
APPLICANT: Lodges, Michael J.
APPLICANT: Lodges, Michael J.
APPLICANT: Sleath, Paul R.
APPLICANT: Houghton, Raymond L.
APPLICANT: Moneill, Particia D.
APPLICANT: Moneill, Particia D.
APPLICANT: Moneil, Mary J.
APPLICANT: Homer, Mary J.
APPLICANT: Homer, Mary J.
APPLICANT: Homer, Mary J.
APPLICANT: Secrist, Heacher
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS
TITLE OF INVENTION: AND TREATMENT OF B. MICROTI INFECTION
FILE REFERENCE: 21012.1426612
CURRENT APPLICATION NUMBER: US/10/294,443
CURRENT PILING DATE: 2002-11-13
NUMBER OF SEQ ID NOS: 243
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NOS: 218
Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20;
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                                                 Indels
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Pred. No. 3e+02;
3; Mismatches 4;
Score 32.5; DB 10;
Pred. No. 3e+02;
3; Mismatches 4;
                                                                                                                                                                                                                                                            Sequence 218, Application US/10294443
Publication No. US20040023865A1
                                                                                               1 LGH-----SDIYSDDKSMKV 15
                                                                                                                          1 LGHHSDMASDINDEEPSFKI 20
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  31.6%;
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Best Local Similarity 40.0%;
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                              8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Babesia microti
US-10-294-443-218
Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea FILE REFERENCE: SPO-103D1
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Sequence 218, Application US/09853079

GENERAL INFORMATION:

APPLICANT: Reed, Steven G.

APPLICANT: Houghton, Raymond L.

APPLICANT: Houghton, Raymond L.

APPLICANT: McNeill, Patricia D.

TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS

TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS

TITLE OF INVENTION: COMPOUNDS AND METHODS

CURRENT APPLICATION NUMBER: US/09/9853,079

CURRENT FILIANT DATE: 201-05-09

NUMBER OF SEQ ID NOS: 224

SEQ ID NO 218

LENGTH: 20
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Pred. No. 1.4;
0; Mismatches 2; Indels
46.6%; Score 48; DB 14; Length 15; 100.0%; Pred. No. 0.64;
                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1). (15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 40
              100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT PILING DATE: 2003-01-29
CURRENT FILING DATE: 2003-01-29
PRIOR APPLICATION NUMBER: PCT/JD97/00740
PRIOR PILING DATE: 1997-03-10
PRIOR PLILING DATE: 1997-03-10
PRIOR PILING DATE: 1998-09-09
NUMBER OF SED ID NOS: 174
SOFTWARE: Patentin version 3.1
LENGTH: 15
                                                                                                                                                                                                                                                          Sequence 54, Application US/10354240 Publication No. US20031185847A1 GENERAL INFORMATION: APPLICANT: Sone, Toshio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           44.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match
Best Local Similarity 80.0.
                                              10; Conservative
                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
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; ORGANISM: Babesia microti
US-09-853-079-218
                                                                                                                        11 KSMKVTVAFN 20
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                   Best Local Similarity
Matches 10; Conserv
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US-09-853-079-218
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  Query Match
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APPLICANT: Deng, Ta
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
FILE REPRENCE: 210121.419C13
CURRENT APPLICATION NUMBER: US/10/079,137B
CURRENT FILING DATE: 2002-02-20
NUMBER OF SEQ ID NOS: 428
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 399
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31.1%; Score 32; DB 15; Length 20; 54.5%; Pred. No. 3.6e+02; i.ve 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 400, Application US/10079137B
; Publication No. US20040073016A1
; GENERAL INFORMATION:
      Sequence 399, Application US/10079137B Publication No. US20040073016A1 GENERAL INFORMATION:
                                                                                                                                                                                                           Wang, Aijun
Skeiky, Yasir A. W.
Harlocker, Susan L.
Day, Craig H.
Li, Samuel X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Fridakis, Tony N. APPLICANT: Reed, Steven G. APPLICANT: Smith, John M. APPLICANT: Misher, Lynda E. APPLICANT: Dillon, Davin C. APPLICANT: Retter, Marc W.
                                                                       APPLICANT: Frudakis, Tony N. APPLICANT: Reed, Steven G. APPLICANT: Smith, John M. APPLICANT: Misher, Lynda E. APPLICANT: Dillon, Davin C. APPLICANT: Retter Marc W.
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Best Local Similarity 54.5
Matches 6; Conservative
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8 HYAIYNEDKLM 18
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US-10-079-137B-400
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ORGANISM: Homo sapiens
US-10-079-137B-399
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US-10-079-137B-400
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                                                                                            US-10-212-679-399

Sequence 399, Application US/10212679

Publication No. US20030125536A1

GENERAL INFORMATION:

APPLICANT: Hirst, Shannon Kathleen

APPLICANT: Hirst, Shannon Kathleen

APPLICANT: Houghton, Ray

APPLICANT: Houghton, Ray

APPLICANT: Houghton, Ray

APPLICANT: Persing, David

APPLICANT: CANCENTION: AND DIAGNOSIS OF BREAST CANCER

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER

FILE REFERENCE: 210121.419C14

CURRENT FILING DATE: 2002-08-02

CURRENT FILING DATE: 2002-08-02

NUMBER OF SEQ ID NOS: 428

SOFTWARE: FastSEQ for Windows Version 4.0

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HENGHI 20
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| Sequence 400, Application US/10212679
| Publication No. US20030125536A1
| GENERAL INFORMATION:
| APPLICANT: Fanger, Gary
| APPLICANT: Hirst, Shannon Kathleen
| APPLICANT: Houghton, Ray
| APPLICANT: Houghton, Ray
| APPLICANT: Kalos, Michael
| TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
| TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
| FILE REFERENCE: 2010.12.4.19014
| CURRENT APPLICATION NUMBER: US/10/212,679
| WUMBER OF SEQ ID NOS: 428
| WUMBER OF SEQ ID NOS: 428
| CORRENT FILLING DATE: 2002-08-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 32; DB 14; Length 20;
Pred. No. 3.6e+02;
2; Mismatches 3; Indels
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Pred. No. 3.6e+02;
2; Mismatches 3
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54.5%;
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Best Local Similarity 54.5
Matches 6, Conservative
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Best Local Similarity 54.5
Matches 6; Conservative
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, ORGANISM: Homo sapiens
US-10-212-679-399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapiens
US-10-212-679-400
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US-10-212-679-400
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US-10-212-679-399
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LENGTH: 20
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APPLICANT: Dillon, Davin C.
APPLICANT: Retter, Marc W:
APPLICANT: Retter, Marc W:
APPLICANT: Wang, Aijun
APPLICANT: Braiky, Vasir A. W.
APPLICANT: Braiky, Vasir A. W.
APPLICANT: Day, Craig H.
APPLICANT: Day, Craig H.
APPLICANT: Deng, Ta
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 21012.419C13
CURRENT APPLICATION NUMBER: US/10/079,137B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
31.1%; Score 32; DB 15; Length 20;
Best Local Similarity 54.5%; Pred. No. 3.6e+02;
Matches 6; Conservative 2; Mismatches 3; Indels
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 400
LENGTH: 20
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RESULT 12

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US-10-033-662-39
US-10-033-662-39
Sequence 39, Application US/10033662
Publication No. US20030092197A1
GENERAL INFORMATION:
TITLE OF INVENTION: Response
TITLE OF INVENTION: Response
TITLE OF INVENTION: Response
TITLE OF INVENTION: Response
TITLE OF INVENTION NUMBER: US/10/033,662
CURRENT APPLICATION NUMBER: US/10/033,662
CURRENT PILING DATE: 2001-12-27
NUMBER OF SEQ ID NOS: 72
SOCTWARE: Patentin version 3.0
SEQ ID NO 39
LENGTH: 14
                                                                            RESULT 14
US-10-033-741-38
US-10-033-741-38
Sequence 38, Application US/10033741
Publication No. US2003009640A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Response
FILE REFERENCE: 9155-079
CURRENT APPLICATION WUMBER: US/10/033,741
CURRENT APPLICATION WUMBER: US/10/033,741
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
SEQ ID NO 38
LENGTH: 14
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29.1%; Score 30; DB 14; Length 14;
Best Local Similarity 42.9%; Pred. No. 5e+02;
Matches 6; Conservative 3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 29.1%; Score 30; DB 14; Length 14; Best Local Similarity 42.9%; Pred. No. 5e+02; Matches 6; Conservative 3; Mismatches 5; Indels
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Job time : 56.5 secs
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1 LGNINTYADDLQNK 14
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3 HYAIYNEDKLM 13
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US-10-033-662-39
                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
GRGANISM: Homo sapiens
US-10-033-741-38
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TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
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ADDRESSEE: ImmuLogic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02154
                                                                                                                                                                                                                                          RESULT 1
US-08-467-023-46
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                                                            January 26, 2005, 15:54:02 ; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                    1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
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3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
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5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-09-142-534D-55
US-08-467-023-234
US-08-467-023-236
US-08-467-023-241
US-08-467-023-241
US-08-467-023-253
US-08-467-023-253
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S-08-467-023-122
S-08-467-023-230
S-08-467-023-257
S-08-467-023-256
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S-08-467-023-247
S-08-467-023-245
S-08-467-023-245
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US-08-467-023-244
US-08-467-023-248
US-08-467-023-250
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                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
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-09-142-524D-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-467-023-46
                                                                                                                                                                                     478139 seqs, 66318000 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                         - protein search, using sw model
                                                                                                                               1 LGHSDIYSDDKSMKVTVAFN 20
                                                                                                                                                   BLOSUM62
Gapop 10.0 , Gapext 0.5
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                                                                                                         US-09-202-464-23
103
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Match Length
          Copyright
                                                                                                                                                                                                                               Minimum DB seq length: 0 Maximum DB seq length: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title:
Perfect score:
                                                                                                                                                    Scoring table:
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                                                                                                                                                                                                                                                                                                           Database :
                                                                                                                                Sequence:
                                                                                                                                                                                     Searched:
                                                             Run on:
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Sequence 47, Appl
Sequence 54, Appl
Sequence 249, Appl
Sequence 60, Appl
Sequence 60, Appl
Sequence 13, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 65, Appl
Sequence 65, Appl
Sequence 65, Appl
Sequence 65, Appl
Sequence 19, Appl
 Sequence
Sequence
Sequence
                          Sequence
Sequence
Sequence
Sequence
US-08-467-023-254
US-08-467-023-251
US-08-467-023-252
US-08-467-023-256
US-08-467-023-236
US-08-467-023-247
US-09-142-524D-54
US-08-945-168-66
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US-08-81-27-946-4
US-08-81-27-946-4
US-08-81-27-946-4
US-08-81-27-946-4
US-08-81-85-86-61
 116
118
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# ALIGNMENTS

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TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING APPLICATION: 424
PRIOR APPLICATION NUMBER: 08/350,225
FILING APPLICATION NUMBER: 08/350,225
FILING APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane R. Remillard
REGISTRATION NUMBER: 38 872
REFERENCE/DOCKET NUMBER: 02.5.6 USD2 (IMI-028CPD2)
TELERPONDICATION INFORMATION:
TELEPRAX: (617) 227-7400
TELEPRAX: (617) 227-7400
TELEPRAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            025.6 USD2 (IMI-028CPD2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: ImmuLogic Pharmaceutical Corporation, STREET: 610 Lincoln St.
Sequence 6, Application US/08467023
Patent No. 6090386
GENERAL INCOMPATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bond, Julian F.;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Garman, Richard D;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Weng, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
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US-09-142-524D-55
JS-09-142-524D-56
                                                                                                                                                                                                                                                                                                                      FEATURE
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                                                           Gaps
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                                                                                                                                                                                                                                         APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bollock, Joanne;
APPLICANT: Bollock, Joanne;
APPLICANT: Garman, Richard D;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Taley, Mark A.;
APPLICANT: Taley, Mark A.;
APPLICANT: Taley, Mark A.;
APPLICANT: Covers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73.8%; Score 76; DB 3; Length 20; 93.8%; Pred. No. 2.3e-06; tive 0; Mismatches 1; Indels
                            91.3%; Score 94; DB 3; Length 20; 90.0%; Pred. No. 2e-09; 1ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 234, Application US/08467023 Patent No. 6090386 GENERAL INFORMATION:
                                                                                           1 LGHSDIYSDDKSMKVTVAFN 20
                                                                                                                1 LGHDDAYSDDKSMKVTVAFN 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DAYSDDKSMKVTVAFN 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: ImmuLogic Pl
STREET: 610 Lincoln St
CITY: Waltham
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 73.8
Best Local Similarity 93.8
Matches 15, Conservative
                            Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-08-467-023-234
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                                                                                                                                                                    RESULT 2
US-08-467-023-234
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    US-08-467-023-46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
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RESULT 3

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APPLICANT: Sone, Toshio
APPLICANT: Boine, Akinori
APPLICANT: Mame, Akinori
APPLICANT: Kino, Kohsuke
APPLICANT: Rino, Kohsuke
TITILE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Wama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Monerale
APPLICANT: Monerale
APPLICANT: Monerale
APPLICANT: Monerale
APPLICANTON: Period Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: 1998-109-09
CURRENT FILING DATE: 1998-109-09
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 55
LENGTH: 15
TYPE: PRT
APPLICANTSM: Cryptomeria japonica
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
67.0%; Score 69; DB 4; Length 15;
Best Local Similarity 86.7%; Pred. No. 2.6e-05;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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; NAME/KEY: MISC FEATURE
; LOCATION: (1)...(15)
; LOCATION: (10...(15)
US-09-142-524D-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: MISC_FEATURE
LOCATION: (1)...(15)
CHER INFORMATION: Cryjl peptide, Figure 1, Row 42
MS-09-142-524D-56
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68.9%; Score 71; DB 4; Le
Best Local Similarity 100.0%; Pred. No. 1.2e-05;
Matches 14; Conservative 0; Mismatches. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 55, Application US/09142524D Patent No. 6719976
Sequence 56, Application US/09142524D Patent No. 6719976
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 LGHDDAYSDDKSMKV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 LGHSDIYSDDKSMKV 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 YSDDKSMKVTVAFN 15
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DATICANT: EXLEY, Mark A.;

TITLE OF INVENTION: Allergenic Proteins And Peptides From
TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SECURIOES: 261
CORRESPONDENCE ADDRESS:
ADDRESSE: Immulogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
STATE: MA
COUNTRY: USA
ZIP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 246, Application US/08467023

Patent No. 6090386

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT:
APPLICANT:
Bond, Julian F.;
APPLICANT:
APP
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORENT APPLICATION DATA:
FILLING DATE: June 6, 1995
CLASSIFICATION A24
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILLING DATE: December 6, 1994
ATTORNEY/AGENT INPORMATION:
MAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 08/350,56 USD2 (IMI-028CPD2)
TELECPHONE: (617) 227-7400
TELECPHONE: (617) 227-7400
TELEPHONE: (617) 227-5941
INPORMATION POR SEQ ID NO: 239:
CENACTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immulogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 60; DB 3;
Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     58.3%; Sc.
100.0%; Pre
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Best Local Similarity 100.
Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 DDKSMKVTVAFN 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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0
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APPLICANT: Bond, Julian F;
APPLICANT: Ruo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H;
APPLICANT: Yeung, Siu-mei H;
APPLICANT: Brauer, Andrew;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Appanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 60; DB 3; Length 17; Pred. No. 0.001; 0; Mismatches 0; Indels
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100.0%; Pred. No. v.
                                                                                                                                                                                                                    Sequence 240, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-467-023-239
; Sequence 239, Application US/08467023
; Patent No. 6090386
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Kuo, Mei-Chang;
Yeung, Siu-mei H.;
Brauer, Andrew;
                                                                                                                                                                                                                                                                                                                                           APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Griffeth, Irwin J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pollock, Joanne;
Bond, Julian F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 58.3
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9 DDKSMKVTVAFN 20
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; FRAGMENT TYPE: internal
US-08-467-023-240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Griffet
APPLICANT: Bond, M
APPLICANT: Garman,
APPLICANT: Garman,
APPLICANT: Kuo, Me:
APPLICANT: Yeung, Ke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ò
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Gaps
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 54.4%; Score 56; DB 3; Length 17; Best Local Similarity 91.7%; Pred. No. 0.0049; Matches 11; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER REALINED FORMY

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Data PC compatible
COMPUTER: Data PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INPORMATION:
NAME: Jane E. Remilland
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 327-7400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-740
TELEPHONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 242:
SEQUENCE CHARACTERESTICS:
LENGTH: 17 aming acids
                         NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPK: (617) 227-5941
INFORMATION FOR SEQ ID NO: 241:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 242, Application US/08467023
Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
     ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 DDKSMKVTVAFN 20
                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Griffe
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USA
                                                                                                                                                                                                                                                                                                                                                                                     US-08-467-023-241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE:
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APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 56; DB 3; Length 16;
Pred. No. 0.0045;
1; Mismatches 0; Indels
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 327-7400
TELEPAX: (617) 227-740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

SOFTWARE: PatentIn Release #1.0, Version #1.25

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/467,023

FILING DATE: June 6, 1995

CLASSIFICATION DATA:

APPLICATION NUMBER: 08/350,225

FILING DATE: December 6, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 241, Application US/08467023 Patent No. 6090386 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
Garman, Richard D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 54.4%;
Best Local Similarity 91.7%;
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 DDKSMKVTVAFN 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-467-023-246
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peptide

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APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Extey, Mark A.;
APPLICANT: Extey, Mark A.;
APPLICANT: Dowers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESSED: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 54.4%; Score 56; DB 3; Length 19; Best Local Similarity 91.7%; Pred. No. 0.0056; Matches 11; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION DATE: US/08/467,023
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 30,5025 (IMI-028CPD2)
TELECOMMUTCATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            025.6 USD2 (IMI-028CPD2)
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Patent No. 6090386
GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Slu-mei H.;
                                                                                      Application US/08467023
                                                                                                                                         GENERAL INFORMATION:
APPLICANT: GRIffeth, Irwin J.;
APPLICANT: POllock, Joanne;
APPLICANT: Bond, Julian F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617) 227-5941
INPORMATION FOR SEQ ID NO: 227:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: MA
COUNTRY: USA
ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 DDKSMKVTVAFN 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-467-023-227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                      ; Sequence 227, Appl. ; Patent No. 6090386
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bollock, Joanne;
APPLICANT: Bollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Faruer, Andrew;
APPLICANT: Exlev, Mark A.;
APPLICANT: Exlev, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Allargenic Corresponding Correspondences: 261
CORRESPONDENCE ADDRESS:
                                                                                                      Query Match

54.4%; Score 56; DB 3; Length 17;
Best Local Similarity 91.7%; Pred. No. 0.0049;
Matches 11; Conservative 1; Mismatches 0; Indels
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Best Local Similarity 91.7%; Pred. No. 0.0052;
Matches 11; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: ImmuLogic Pharmaceutical Corporation, Inc. 610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: June 6, 1995
CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane B. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-7400
TELEFAX: (617) 227-75941
INPORMATION FOR SEQ ID NO: 253:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: "".

COUNTRY: USA
ZID: 02154
ZID: 02154
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DoS/MS-DOS
SOFTWARE: ParentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
TITING DATE: Unne 6, 1995
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 253, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 amino acids
                                  internal
                                                                                                                                                                                                               9 DDKSMKVTVAFN 20
                                                                                                                                                                                                                                          9 DDKSMKVTVAFN 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Imm.
STREET: 610 Lir
CITY: Waltham
STATE: MA
; MOLECULE TYPE:
; FRAGMENT TYPE:
US-08-467-023-242
                                                                                                                                                                                                                                                                                                                                                                     US-08-467-023-253
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ratent No. 6090386

GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Prauer, Andrew;
APPLICANT: Prauer, Andrew;
APPLICANT: Brauer, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50.5%; Score 52; DB 3; Length 19; 83.3%; Pred, No. 0.027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc. STREET: 610 Lincoln St CITY: Waltham
              COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPAX: (617) 227-7400
TELEFAX: (617) 227-7401
TELEPAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
TVVPE: maino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR PALICATION DATA:
APPLICATION NUMBER: 08/350,225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
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; Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 50.5
Best Local Similarity 83.3
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-08-467-023-122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 DDKSMKVTVAFN 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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US-08-467-023-230
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; Sequence 122, Application US/08467023
; Patent No. 609386
; GENERAL INFORMATION:
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Bond, Julian F.;
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS: 261
CORRESPONDENCE ADDRESS: ImmiLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Stewart, Stewart Promers, Stewart Promers, Stewart Promers, Stewart Powers, P
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pred. No. 0.027;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                     STREET: Immunogic Fnarmaceurical Corporation, inc.
STATEET: Waltham
STATE: WA
COUNTRY: Waltham
CONTUTRY: WAS
ZIP: 02154
COMPUTER: BADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ELADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: DATE FLOPEN BADA:
COMPUTER: DATE FLOPEN BADA:
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
FILING DATE: Unne 6, 1995
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKT NUMBER: 38,872
REFERENCE/DOCKT NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPAN: (617) 227-7400
TELEFAX: (617) 227-7400
TELEFAX: (617) 227-7400
TELEFAX: (617) 227-7401
                                                                                                                                                                                                                                                                                                                             B: ImmuLogic Pharmaceutical Corporation, Inc.
610 Lincoln St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 83.3%;
Matches 10; Conservative
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; FRAGMENT TYPE: internal
US-08-467-023-121
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GY: linear
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TOPOLOGY: linear

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APPLICANT: Griffeth, Irwin J.;

APPLICANT: Carifeth, Jonne;

APPLICANT: Pollock, Joanne;

APPLICANT: Bond, Julian F.;

APPLICANT: Graman, Richard D;

APPLICANT: Graman, Richard D;

APPLICANT: Earley, Mark A.;

APPLICANT: Exley, Mark A.;

APPLICANT: Mark A.;

COUNTRY: Waltham

STAFE: Ma.

COMPUTER: INPECTION DATA:

COMPUTER: PatentION DATA:

COMPUTER: PatentION DATA:

COMPUTER: PatentION DATA:

APPLICATION NUMBER: Us/08/467,023

FILING DATE: December 6, 1994

ATTORNEY AGENT INPORMATION:

APPLICATION NUMBER: 025.6 USD2 (IMI-028CPD2)

TELEPHONE: Jan MONDER: 327-5941

INPORMATION FOR EXQ IN NO: 257:

EELECOMMUTER: TRANDER SEQ ID NO: 257:

EELECOMMUTER SEQUENCE CAREAGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
                                                                                                                                                                                                                                                                                                                                                                                Score 52; DB 3; Length 20;
Pred. No. 0.028;
2; Mismatches 0; Indels
FILING DATE: December 6, 1994

ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38, 872

REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-5940

INFORMATION FOR SEQ ID NO: 230:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 257, Application US/08467023 Patent No. 6090386
                                                                                                                                                                                                                                                                                                                                                                                Query Match 50.5%;
Best Local Similarity 83.3%;
Matches 10; Conservative
                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
J: FRAGMENT TYPE: internal
US-08-467-023-230
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EEKSMKVTVAFN 13
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i MOLECULE TYPE: peptide
j; FRAGMENT TYPE: internal
US-08-467-023-257
Query Match

Query Match
Matches 10; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
Qy 9 DDKSMKYTVAFN 20
::||||||||
Db 1 NNKSMKYTVAFN 12
Search completed: January 26, 2005, 16:08:27
Job time: 15.6 secs
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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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| Cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/DSUG_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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| Cgn2_6/ptodata/1/pubpaa/US096_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US096_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*
| CGn2_6/ptodata/1/pubpaa/US108
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/cgn2_6/ptodata/1/pubpaa/US60_PUBGOMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1608061 seqs, 361289386 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Published_Applications_AA:*
                                                                                                                                                                     OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-202-464-24
103
1 KSMKVTVAFNQFGPNAGQRM 20
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 20
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                             Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

					SUMMARIES	
		<b>æ</b> (				
No.	Score	Match	Query Match Length DB	DB	ID	Description
	78	75.7	15	14	US-10-354-240-9	Sequence 9. Appli
7	7.8	75.7	15	14	US-10-354-240-57	Sequence 57, Appl
3	78	75.7	15	14	US-10-354-240-158	158
4	94	73.8	15	14	US-10-354-240-58	
2	51	49.5	15	14	US-10-354-240-59	59,
9	48	46.6	15	14	US-10-354-240-56	56,
7	30	29.1	o	14	US-10-079-167-73	73,
60	30	29.1	6	16	US-10-653-624-73	73,
თ	30	29.1	თ	16	US-10-833-439-73	73,
10	30	29.1	σ	17	US-10-833-745-73	
11	30	29.1	6	17	US-10-833-744-73	73,
12	30	29.1	o	17	US-10-686-943-73	73,
13	29.5	28.6	20	14	US-10-282-960-74	74

	Sequence 1288, Ap Sequence 1508, Ap Sequence 8, Appli Sequence 8, Appli Sequence 8, Appli	<b>ຜ</b> ິສິສິສິສິ	33, 19, 19, 19, 19, 19, 19, 19, 19, 19, 19	Sequence 114, App Sequence 8, Appli Sequence 29, Appl Sequence 29, Appl Sequence 4176, Ap Sequence 13, Appl Sequence 2629, Ap Sequence 2629, Ap Sequence 2629, Ap
4444	14 US-10-225-567A-1288 14 US-10-225-567A-1508 US-09-776-874A-8 9 US-09-988-113-8 14 US-10-341-582-8	US-10-384- US-10-384- US-10-371- US-10-456- US-10-785-	17 US-10-745-242A-33 16 US-10-203-915A-34 16 US-10-203-915A-35 9 US-09-826-752-19 17 US-10-729-441-37 17 US-10-729-441-38 18 US-08-841-636A-29	14 US-10-300-694A-114 US-10-76-8139A-8 16 US-10-825-978-29 16 US-10-149-138-4176 15 US-10-132-284A-13 17 US-11-149-138-2629 15 US-10-149-138-2629 15 US-10-149-138-2629
9 9 9 8 8	<b></b>			99999999
2 2 2 2 2 2 2 2 2 2 2 3 3 3 3 3 3 3 3 3	28. 27. 27.			27. 27. 27. 26. 26. 26.
	19 20 21 22 23 23			0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0

## ALIGNMENTS

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TUBLICANTS SONE, TOSALO

APPLICANT: SONE, Toshio

APPLICANT: SONE, Toshio

APPLICANT: Sone, Toshio

APPLICANT: Kume, Akinori

APPLICANT: Tume, Akinori

APPLICANT: Time, Missue

APPLICANT: Time, Missue

APPLICANT: Time, Missue

TILES FERRENCE: SPO-103D1

CURRENT APPLICATION NUMBER: US/10/354,240

CURRENT APPLICATION NUMBER: PCT/JP97/00740

PRIOR APPLICATION NUMBER: US 09/142,524

PRIOR APPLICATION NUMBER: US 09/142,524

PRIOR APPLICATION NUMBER: US 09/142,524

NUMBER: OF SEQ ID NOS: 174

SEQ ID NOS: 174

SEQ ID NOS: 174
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0
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Pred. No. 1.9e-06;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75.7%; Scor
100.0%; Pre
Sequence 9, Application US/10354240 Publication No. US20030185847A1
                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Cryptomeria japonica
US-10-354-240-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KSMKVTVAFNQFGPN 15
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Best Local Similarity 100.
Matches 15; Conservative
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RESULT

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APPLICANT: Cume, Akinori
APPLICANT: Cume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Lairiki, Cohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: UNMBER: US/10/354,240
CURRENT FILING DATE: 2003-01-29
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR APPLICATION NUMBER: US 09/142,524
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SUSTIMER: Patentin version 3.1
SEQ ID NO 58
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Wama, Akinori
APPLICANT: Wama, Akinori
APPLICANT: Wino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Diseas
FILE REPERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT FILING DATE: 1997-03-10
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SEQ ID NOS: 174
SEQ ID NOS: 174
LENGTH: 15
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; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 45
US-10-354-240-59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; LOCATION: (1)...(15); OTHER INFORMATION: Cryjl peptide, Figure 1, Row 44 US-10-354-240-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73.8%; Score 76; DB 14;
93.3%; Pred. No. 4.2e-06;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 59, Application US/10354240; Publication No. US20030185847A1; GENERAL INFORMATION:
                                                                                                                                                    Sequence 58, Application US/10354240
Publication No. US20030185847A1
GENERAL INFORMATION:
APPLICANT: Sone, Toshio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Cryptomeria japonica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 TVAFNOFGPNAGORM 20
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Query Match
Best Local Similarity 93.3%
Matches 14; Conservative
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NAME/KEY: MISC_FEATURE
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Publication No. US20030185847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Name, Akinori

APPLICANT: Name, Akino

APPLICANT: Name, Akino

APPLICANT: Name, Akino

APPLICANT: Name, Akino

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

CURRENT APPLICATION NUMBER: US/10/354,240

CURRENT APPLICATION NUMBER: DET/JP97/00740

PRIOR APPLICATION NUMBER: DET/JP97/00740

PRIOR APPLICATION NUMBER: US 09/142,524

PRIOR PILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFTWARE: PatentIn version 3.1
                                                                                APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Sone, Toshio
APPLICANT: Mame, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION NUMBER: US/10/354,240
CURRENT APPLICATION NUMBER: PCT/JP97/00740
PRIOR PILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOGTWARE: Patentin version 3.1
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hes 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; LOCATION: (1)..(15); OTHER INFORMATION: Cryjl peptide, Figure 1, Row 43 US-10-354-240-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 75.7%; Score 78; DB 14; I Best Local Similarity 100.0%; Pred. No. 1.9e-06; Matches 15; Conservative 0; Mismatches 0;
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OTHER INFORMATION: Figure 7, Row a
                          Sequence 57, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
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LENGTH: 15
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LENGTH: 15
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1 KSMKVTVAFNQFGPN 15

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APPLICANT: Gilbert, Sarah C.
APPLICANT: Schneider, Jorg
APPLICANT: Plebanski, Magdalena
APPLICANT: Hanke, Tomas
APPLICANT: Hanke, Tomas
APPLICANT: Blanchard, Tom
APPLICANT: Smith, Geoffrey L.
APPLICANT: Blanchard, Tom
TITLE OF INVENTION: Methods and Reagents for Vaccination
TITLE OF INVENTION: Mothods and Reagents for Vaccination
TITLE OF INVENTION: Mothods and Reagents for Vaccination
TITLE OF INVENTION: Mothods and Reagents for Vaccination
TITLE OF INVENTION: MOTHORER: US/10/653,624
CURRENT APPLICATION NUMBER: US/09/454,204A
PRIOR FILING DATE: 1999-12-09
PRIOR FILING DATE: 1998-6-09
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FASISEQ for Windows Version 4.0
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                                                                                                                      FEATURE:
; OTHER INFORMATION: CTL Peptide Epitope of P. falciparum TRAP
US-10-079-167-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: CTL Peptide Epitope of P. falciparum TRAP US-10-653-624-73
                                                                                                                                                                                                                            Length 9;
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                                                                                                                                                                                                                       Score 30; DB 14;
Pred. No. 1.5e+06;
2; Mismatches 1;
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Pred. No. 1.5e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Mismatches
     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 73 LENGTH: 9
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. Sequence 73. Application US/10833439
. Publication No. US20040175365A1
. GENERAL INFORMATION:
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29.1%; Scc
Best Local Similarity 62.5%; Pre
Matches 5; Conservative 2;
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Plebanski, Magdalena
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Best Local Similarity 62.5%;
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Smith, Geoffrey L.
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1 INVAFNRF 8
                                                                        TYPE: PRT
ORGANISM: Unknown
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i Sequence 56, Application US/10354240

i Publication No. US20030185847A1

i GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Name, Akinori

APPLICANT: Name, Akinori

APPLICANT: Name, Akinori

APPLICANT: Nico, Kobauke

ITILE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

FILE REFERENCE: SPO-103D1

CURRENT PILING DATE: 2003-01-29

PRIOR APPLICATION NUMBER: PCT/JP97/00740

PRIOR PELICATION NUMBER: US 09/142,524

PRIOR PILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SEQ ID NO 56

LENGTH: 15
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46.6%; Score 48; DB 14; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.29;
Matches 10; Conservative 0; Mismatches 0; Indels
Query Match
49.5%; Score 51; DB 14; Length 15;
Best Local Similarity 90.0%; Pred. No. 0.089;
Matches 9; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: MISC_FEATURE

1 LOCATION: (1)..(15)

2 OTHER INFORMATION: Cryjl peptide, Figure 1, Row 42

US-10-354-240-56
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GENERAL INFORMATION:

APPLICANT: Hill, Adrian V.S.

APPLICANT: Gilbert, Sarah C.

APPLICANT: Gilbert, Sarah C.

APPLICANT: Gilbert, Sarah C.

APPLICANT: Rece, William

APPLICANT: Schneider, Joerg

TITLE OF INVENTION: Vaccination Method

FILE REFERENCE: 2907.1000-001

CURRENT APPLICATION NUMBER: US/10/079,167

CURRENT PILING DATE: 1999-12-09

PRIOR PILING DATE: 1999-12-09

PRIOR PILING DATE: 1999-06-09

PRIOR PILING DATE: 1999-06-09

PRIOR PILING DATE: 1999-06-09

PRIOR PILING DATE: 1997-06-09

PRIOR PILING DATE: 2001-09-13

PRIOR PILING DATE: 2001-09-13

PRIOR PILING DATE: 2001-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Cryptomeria japonica
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                                                                                                                                            1 OFGPNCGORM 10
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1 INVAFNRF
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Publication No. US20040191272A1

GENERAL INFORMATION:
APPLICANT: Mill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Schneider, Joza and Reagents for Vaccination
TITLE OF INVENTION: Methods and Reagents for Vaccination
TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response
FILE REFERENCE: 2907.100.000
CURRENT APPLICATION NUMBER: US/10/686,943
FRIOR FILING DATE: 2003-10-16
FRIOR FILING DATE: 1999-06-09
FRIOR FILING DATE: 1999-06-09
FRIOR FILING DATE: 1998-06-09
FRIOR FILING DATE: 1998-06-09
FRIOR FILING DATE: 1998-06-09
NUMBER OF SEQ ID NOS: 78

SEQ ID NO 7: BESTERE FASTERE 
## APPLICANT: Blanchard, Tom
### TITLE OF INVENTION: Methods and Reagents for Vaccination
### TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response
### FILE OF INVENTION: Which Generate A CD8 T Cell Immune Response
### FILE REFERENCE: 2907.1000-000
### CURRENT APPLICATION NUMBER: US/10/686,943
### PRIOR APPLICATION NUMBER: US/10/686,943
### PRIOR PILING DATE: 2003-10-16
### PRIOR PILING DATE: 1999-12-09
### PRIOR FILING DATE: 1999-12-09
### PRIOR PRIOR DATE: 1999-12-09
### PRIOR P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: CTL Peptide Epitope of P. falciparum TRAP
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29.1%; Score 30; DB 17; Length 9;
Best Local Similarity 62.5%; Pred. No. 1.5e+06;
Matches 5; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29.1%; Score 30; DB 16; Length 9; 62.5%; Pred. No. 1.5e+06; tive 2; Mismatches 1; Indels
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Matches 5; Conservative
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1 INVAFNRF 8
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ORGANISM: Unknown
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US-10-833-745-73
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                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 32, Application US/10730454
; Sequence 32, Application US/10730454
; Publication No. US200401757571
; RENERAL INFORMATION:
; APPLICANT: Olsen, Arne
; APPLICANT: Roggen, Erwin
; APPLICANT: Ernet, Steffen
; TITLE OF INVENTION: Low Allergenic Protein Variants
; FILE REFERENCE: 5676.200-US
; CURRENT APPLICATION NUMBER: US/09/417,608
; PRIOR FILING DATE: 1999-10-13
                                                                                                              28.2%; Score 29; DB 16; I 100.0%; Pred. No. 1.5e+06; tive 0; Mismatches 0;
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE: OTHER INFORMATION: Synthetic
             OTHER INFORMATION: Synthetic
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Matches 5; Conservative
                                                                                                              Query Match 28.2
Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                             13 GPNAG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chen, Si-Yi
APPLICANT: Chen, Si-Yi
APPLICANT: Chen, Si-Yi
APPLICANT: Chen, Si-Yi
APPLICANT: Chen, Si-Xi
APPLICANT: Schroers, Roland
APPLICANT: Schroers, Roland
TITLE OF INVENTION: Associated Antigen
FILE REFERENCE: P02193USI
CURRENT APPLICATION NUMBER: US/10/282,960
CURRENT APPLICATION NUMBER: US 60/345,012
PRIOR APPLICATION NUMBER: US 60/345,012
PRIOR PILING DATE: 2001-10-29
NUMBER OF SEQ ID NOS: 100
SOFTWARE: Patentin version 3.1
SEQ ID NO 74
LENGTH: 20
ILENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                            Query Match
29.1%; Score 30; DB 17; Length 9;
Best Local Similarity 62.5%; Pred. No. 1.5e+06;
Matches 5; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 26, Application US/10730454
; Bublication No. US2004017575A1
; GENERAL INFORMATION:
; APPLICANT: Olsen, Arne
; APPLICANT: Ernst, Steffen
; TITLE OF INVENTION: Low Allergenic Protein Variants
; FILE REFERENCE: 56.200-US
; CURRENT FILING DATE: 2003-12-08
; PRIOR APPLICATION NUMBER: US/10/730,454
; PRIOR APPLICATION NUMBER: US/9/417,608
; PRIOR PRINING DATE: 1999-10-13
; NUMBER OF SEQ ID NOS: 101
; SEQ ID NO 26
; LENGTH: 9
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 73
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 SMKVTVAFNQFGPNAGORM 20
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1 INVAFNRF 8
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ORGANISM: Artificial
                                                                                                                                                                        TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 13
US-10-282-960-74
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US-10-730-454-26
                                                                                                                                                                                                                                       FEATURE:
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Gaps

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Sequence 47, Appl
                                                           January 26, 2005, 15:54:02; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Description
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Sequence
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Sequence
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-467-023-230
US-09-142-524D-58
US-08-467-023-121
US-08-467-023-122
US-08-467-023-126
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US-09-142-524D-158
US-08-467-023-227
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-08-467-023-235
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-08-467-023-256
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                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                          - protein search, using sw model
                                                                                                        US-09-202-464-24
103
1 KSMKVTVAFNQFGPNAGQRM 20
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Gapop 10.0 , Gapext 0.5
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Match Length DB
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                                                                                                        Title:
Perfect score:
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                                                               Run on:
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No.
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```

## ALIGNMENTS

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US-08-47-023-47

US-08-47-023-47

US-08-47-023-47

Sequence 47, Application US/08467023

Fatent No. 6090386

GENERAL INFORMATION

APPLICANT: Griffeth, Irwin J.;

APPLICANT: Garman, Richard D;

APPLICANT: Garman, Richard D;

APPLICANT: Raugh, Mai-Chang;

APPLICANT: Raugh, Mai-Chang;

APPLICANT: Raugh, Mai-Chang;

APPLICANT: Raugh, Mai-Chang;

APPLICANT: Powers, Steven P.

ITILE OF INVENTION: Japanese Cedar Pollen

NUMBER OF SEQUENCES: 26

CONDERSES: ImmuLogic Pharmaccutical Corporation, Inc.

STREET: Gio Lincoln St.

COUNTRY: USA

COUNTRY: USA

ZIPR: MA

COUNTRY: USA

ZIPR: Patentin Release #1.0, Version #1.25

COUNTRY: USA

ZIPR: Patentin Release #1.0, Version #1.25

COUNTRY: USA

ZIPR: Date Date Date OFFERTATION NUMBER: US(8)/467, 023

FILING DATE: Unne 6, 1995

CLASSIFICATION NUMBER: US/8150

FILING DATE: December 6, 1994

APPLICATION NUMBER: 08/350,225

FILING DATE: December 6, 1994

APPLICATION NUMBER: 38, 972

RESERRENCE/DOCKEY NUMBER: 38, 972

RESERVENCE/DOCKEY ```

```
GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Mame, Akinori
APPLICANT: Lwama, Akinori
APPLICANT: Iwama, Akiko
TITLE OF INVENTION NUMBER: 1990-103
CURRENT FILING DATE: 1997-03-10
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 158
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                                  Gaps
  Gaps
  .;
0
   APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne,
APPLICANT: Bond, Julian F.;
APPLICANT: Bornd, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Allergenic Correspondences: 261
CORRESPONDENCE ADDRESS:
                                    ;
0
   Length 15;
  Indels
   B: ImmuLogic Pharmaceutical Corporation, Inc.
610 Lincoln St
                                    Indels
                Pred. No. 2.2e-07;
Mismatches 0;
  .2e-07;
   DB 4;
   0; Mismatches
   75.7%; Score 78; 100.0%; Pred. No.
   Sequence 158, Application US/09142524D Patent No. 6719976
   ; Sequence 227, Application US/08467023
; Patent No. 6090386
; GENERAL INFORMATION:
; APPLICANT: Griffeth, Irwin J.;
   E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
  , LOCATION: (1). (15)
, OTHER INFORMATION: Figure 7, Row a
US-09-142-524D-158
100.0%; FIT
   TYPE: PRT
ORGANISM: Cryptomeria japonica
   1 KSMKVTVAFNQFGPN 15
  1 KSMKVTVAFNOFGPN 15
  1 KSMKVTVAFNQFGPN 15
                                       15; Conservative
   Query Match 75.7
Best Local Similarity 100.
Matches 15; Conservative
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  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
   FEATURE:
NAME/KEY: MISC_FEATURE
   COMPUTER: IBM PC
OPERATING SYSTEM:
                  Best Local Similarity
   Waltham
   USA
  ADDRESSEE:
   US-09-142-524D-158
  US-08-467-023-227
   COUNTRY:
   STREET:
   Matches
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   ð
  APPLICANT: Sone, Toshio
APPLICANT: Sume, Akinori
APPLICANT: Mane, Akinori
APPLICANT: Mane, Akinori
APPLICANT: Mane, Akinori
APPLICANT: Mane, Akivo
APPLICANT: Mane, Akivo
APPLICANT: Wama, Akivo
APPLICANT: Wana, Akivo
APPLICANT: Wana, Akivo
APPLICANT: Wana, Akivo
APPLICANT: Mane, Mobile
FILE REFERENCE: SPO-103
FILE REFERENCE: SPO-103
FILE REFERENCE: SPO-103
FILE REFERENCE: 1994-09
FRIOR APPLICATION NUMBER: PCT/JP97/00740
FRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 57
FILENGTH: 15
FILENGTH: 15
FILENGTH: 15
   APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Twine, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Trino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: PCT/0P97/00740
PRIOR APPLICATION NUMBER: PCT/0P97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 9
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  Query Match 96.1%; Score 99; DB 3; Length 20; Best Local Similarity 95.0%; Pred. No. 5.4e-11; Matches 19; Conservative 0; Mismatches 1; Indels
   0; Indels
   LOCATION: (1).7(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 43
   Score 78; DB 4; Li
Pred. No. 2.2e-07;
   75.7%; Scor.
100.0%; Pred. No. ...
  Sequence 57, Application US/09142524D Patent No. 6719976
  Sequence 9, Application US/09142524D
Patent No. 6719976
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   ORGANISM: Cryptomeria japonica
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  Query Match
Best Local Similarity 100.0
Matches 15; Conservative
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  US-09-142-524D-57
  US-09-142-524D-9
   US-09-142-524D-9
                       US-08-467-023-47
  LENGTH
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DB 4; Length 15;

75.7%; Score 78;

Query Match

Gaps

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0
  APPLICANT: Garman, Richard D,
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Yeung, Sudreni H.;
APPLICANT: Yeung, Sudreni H.;
APPLICANT: Braler, Andrew;
APPLICANT: Braler, Mark A.;
APPLICANT: Braler, Steven P.
TITLE OF INVENTION: Japanese Cedar Pollen
CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
  Length 19;
   0; Indels
  STREET: WALTHAM
COTHY: WALTHAM
COTHY: WA
   MEDIUM TYPE: Floppy disk
COMPUTER: ELOppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
FILING DATE: Dacember 6, 1994
PRIOR APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INPORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 0.25-7400
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7401
TELEPHONE: (617) 227-7401
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
  Score 78; DB 3; L. Pred. No. 2.9e-07; 0; Mismatches 0;
   ; Sequence 230, Application US/08467023
; Patent No. 6090386
   GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
  Query Match

Best Local Similarity 100.0%; P
Matches 15; Conservative 0;
  TELEFAX: (617) 227-5941
INPORMATION FOR SEQ ID NO: 236:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
  1 KSMKVTVAFNQFGPN 15
  3 KSMKVTVAFNOFGPN 17
  LENGTH: 20 amino acids
TYPE: amino acid
TOPOLOGY: linear
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; FRAGMENT TYPE: internal
US-08-467-023-236
   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
  COUNTRY: USA
ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
   amino acid
   RESULT 7
  ò
   셤
  ö
  Gaps
  ö
   APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Bond, Julian F.;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Bxley, Mark A.;
APPLICANT: Joannese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSE: Immut.~~¹
STRREY
STRREY: Immut.~~¹
   Score 78; DB 3; Length 19;
Pred. No. 2.9e-07;
   0; Indels
  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
FILING DATE: June 6, 1995
FILING APPLICATION: 424
PRIOR APPLICATION: DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERRINGE/POCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 227.760
SEQUENCE CHARACTERISTICS:
LENGTH: 19 main acide
   COUNTRY: USA
ZIP: 02154
COUNTRY EADABLE FORM:
MEDUTER READABLE FORM:
MEDUTER TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: Unne 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
  AFFIREATION NUMBERS 07320,622

PILING DATE: December 6, 1994

ATTORNEY/AGENT INFORMATION:

NAME: Jane E. Remillard

NUMES: JANE E. Remillard

REFERENCE/DOCKET NUMBER: 38, 872

REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
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   3 KŚMKVTVAFNOFGPN 17
   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
  TYPE: amino acid
  US-08-467-023-227
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   Proteins And Peptides From
  Score 74; DB 3; Length 19;
Pred. No. 1.5e-06;
   1; Indels
   ImmuLogic Pharmaceutical Corporation, Inc
                 URREATION SISTEM:

URREAT APPLICATION DATA:

CURREAT APPLICATION DATA:

APPLICATION NUMBER:

FILING DATE:

APPLICATION NUMBER:

CLASSIFICATION NUMBER:

PRIOR APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

ATTORNEY/AGENT INPORMATION:

NAME:

REGISTRATION NUMBER:

REFERENCE/DOCKET NUMBER:

TELEFHONE:

TELEFHONE:

(617) 227-7400

TELEFRAM:

INPORMATION FOR SEQ ID NO: 121:

SEQUENCE CHARACTERISTICS:
  025.6 USD2 (IMI-028CPD2)
  COMPUTER: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: Unne 6, 1995
CIASSIFICATION: 424
  TITLE OF INVENTION: Allergenic Proteins Ar
TITLE OF INVENTION: Japanese Cedar Pollen
   0; Mismatches
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
   Sequence 122, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
    PC-DOS/MS-DOS
   Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
Garman, Richard D;
Kuo, Mei-Chang;
  Yeung, Siu-mei H.;
Brauer, Andrew;
Exley, Mark A.;
Powers, Steven P.
  Query Match 71.8%;
Best Local Similarity 93.3%;
Matches 14; Conservative
  1 KSMKVTVAFNOFGPN 15
   3 KSMKATVAFNQFGPN 17
  STREET: 610 Lincoln St
   19 amino acids
  NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immulogic
   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
  linear
        OPERATING SYSTEM:
  amino acid
   FILING DATE: Ju:
CLASSIFICATION:
  Waltham
   USA
  ZIP: 02154
  APPLICANT:
APPLICANT:
APPLICANT:
  APPLICANT:
APPLICANT:
APPLICANT:
  US-08-467-023-121
  TOPOLOGY:
   APPLICANT:
   COUNTRY:
  APPLICANT:
   LENGTH:
  STATE:
  ò
   셤
   uairiki, Kazuo

APPLICANT: Iwama, Akiko

APPLICANT: Iwama, Akiko

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

FILE REFRENCE: SPO-103

CURRENT APPLICATION NUMBER: US/09/142,524D

CURRENT PILING DATE: 1998-09-09

PRIOR APPLICATION NUMBER: PCT/JP97/00740

NUMBER OF SEQ ID NOS: 174

SOFTWARE: Patentin version 3.1

LENOTH: 15

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   Gaps
   Sequence 121, Application US/08467023
; Sequence 121, Application US/08467023
; Patent No. 6090386
; GENERAL INFORMATION:
    APPLICANT: Griffeth, Irwin J.;
    APPLICANT: Pollock, Joanne;
    APPLICANT: Garman, Richard D;
    APPLICANT: Yeung, Siu-mei H.;
    APPLICANT: Yeung, Siu-mei H.;
    APPLICANT: Yeung, Siu-mei H.;
    APPLICANT: Exley, Mark A.;
    APPLICANT: Exley, Mark A.;
    APPLICANT: Exley, Mark A.;
    APPLICANT: Powers, Steven P.
    TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
    NUMBER OF SEQUENCES: 261
    CORRESPONDENCE ADDRESS:
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   Query Match
Best Local Similarity 93.3%; Pred. No. 5e-07;
Matches 14; Conservative 0; Mismatches 1; Indels
            Length 20;
   0; Indels
  ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
  , LOCATION: (1).7(15)
, OTHER INFORMATION: Cryjl peptide, Figure 1, Row 44
US-09-142-524D-58
            Score 78; DB 3;
Pred. No. 3e-07;
0; Mismatches 0
75.7%; Scc...
100.0%; Pre
   ZIP: 02154
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
  ORGANISM: Cryptomeria japonica
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Best Local Similarity 100.
Matches 15; Conservative
  NAME/KEY: MISC FEATURE
   STATE: M. COUNTRY:
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RESULT 13
US-08-467-023-235
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  Sequence 126, Application US/08467023

Sequence 126, Application US/08467023

Patent No. 6090386

GENERAL INFORMATION:

APPLICANT: Griffeth, Irwin J.;

APPLICANT: Bollock, Joanne;

APPLICANT: Garman, Richard D;

APPLICANT: Garman, Richard D;

APPLICANT: Mone, Chang;

APPLICANT: Mone, Chang;

APPLICANT: Mone, Situme! H.;

APPLICANT: Andrew;

APPLICANT: Allersenic Proteins And Peptides From

TITLE OF INVENTION: Japanese Cedar Pollen

CORRESPONDENCE ADDRESS:

ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.

GITTY: Waltham

STATE: MA
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   Query Match 71.8%; Score 74; DB 3; Length 19; Best Local Similarity 93.3%; Pred. No. 1.5e-06; Matches 14; Conservative 0; Mismatches 1; Indels
   CURRENT APPLICATION DATA:

SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: US/08/467,025
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,72
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECHONE: (617) 227-7400
TELEPHONE: (617) 227-740
TELEPHONE: (617) 227-7591
INFORMATION FOR SEQ ID NO: 126:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acide
TYPE: amino acide
TYPE: amino acide
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
   COMPUTER READABLE FORM:
BODIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
TYPE: amino acids
   1 KSMKVTVAFNQFGPN 15
   3 KSMKVTAAFNOFGPN 17
   ## MOLECULE TYPE: linear
### MOLECULE TYPE: peptide
### FRAGMENT TYPE: internal
US-08-467-023-122
  MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
   CITY: Walth
STATE: MA
COUNTRY: US
ZIP: 02154
   RESULT 11
US-08-467-023-126
   US-08-467-023-126
  ò
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  ö
  Gaps
  ö
  ;
  GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
68.0%; Score 70; DB 3; Length 19; 86.7%; Pred. No. 7.7e-06; Live 0; Mismatches 2; Indels
  Length 17;
  1; Indels
   CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc. STREET: 610 Lincoln St CITY: Waltham
  PRIOR APPLICATION 1824
PRIOR APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INPORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,072
TELEPHONE: (617) 227-7400:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: US/08/467,023
FILING DATE: Unne 6, 1995
CLASSIFICATION: 424
   Query Match 67.0%; Score 69; DB 3;
Best Local Similarity 93.3%; Pred. No. 1e-05;
Matches 14; Conservative 0; Mismatches
  ; Sequence 257, Application US/08467023
; Patent No. 6090386
  1 KSMKVTVAFNQFGPN 15
  3 KSMKATAAFNOFGPN 17
  1 KSMKVTVAFNQFGPN 15
  TELBFAX: (617) 27-5941
INFORMATION FOR SEQ 1D NO: SEQUENCE CHARACTERISTICS: LENGTH: 17 amino acids
  13; Conservative
   ) MOLECULE TYPE: peptide
) FRAGMENT TYPE: internal
US-08-467-023-257
Query Match
Best Local Similarity
Matches 13; Conserva
  amino acid
  02154
   RESULT 12
US-08-467-023-257
  STATE: M
COUNTRY:
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TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
   ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc. STREET: 610 Lincoln St CITY: Waltham
  ImmuLogic Pharmaceutical Corporation, Inc.
   MEDIUM TYPE: Floppy disk
COMPUTER: ELOMPY disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: Unne 6, 1995
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
  63.1%; Score 65; DB 3; L6 100.0%; Pred. No. 4.6e-05; tive 0; Mismatches 0;
   ; Sequence 256, Application US/08467023
; Patent No. 6090386
; GENERAL INFORMATION:
  APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
   TELEPRAK: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION POR SEQ ID NO: 255
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
  COUNTRY: USA
ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
  1 KSMKVTVAFNQFG 13
   2 KSMKVTVAFNOFG 14
  610 Lincoln St
   Query Match 63.1
Best Local Similarity 100.
Matches 13; Conservative
  MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-467-023-255
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  amino acid
   ADDAL.
STREET: 61.
   MA
USA
   02154
  ADDRESSEE:
   US-08-467-023-256
   STATE: MA
  TOPOLOGY:
  ð
   ö
   Gaps
   ..
0
   GENERAL INFORMATION:
APPLICANT: Giffeth, Irwin J.;
APPLICANT: Golfeth, Irwin J.;
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Fowers, Steven P.
TITLE OF INVENTION: Japanese Cedar Pollen
INVERS OF SEQUENCES: 261
APPLICANT: Fowers, Steven P.
TITLE OF INVENTION: Japanese Cedar Pollen
INVERS OF SEQUENCES: 25
ADDRESSEE: Immulogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Maltham
COUNTR: Maltham
COUNTR: Maltham
COUNTR: Farabable FORM:
MEDIUM TYPE: 10 Lincoln St
CONFUTER: IBM PC Compatible
OPERATING SYSTEM: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: IDM PC Compatible
OPERATING SYSTEM: Ploppy disk
COMPUTER: June 6, 1995
STATE: MA
APPLICATION NUMBER: US/08/467,023
FILING DATE: December 6, 1994
ATTORNEY AGENT INFORMATION:
TELEPHONE: Gill 227-5941
INFORMATION POR SEQ ID NO: 225-5941
INFORMATION POR SEQ ID NO: 235:
SEGUENCE CHARACTERISTICS:
INFORMATION POR SEQ ID NO: 235:
   Query Match 63.1%; Score 65; DB 3; Length 13; Best Local Similarity 100.0%; Pred. No. 3.9e-05; Matches 13; Conservative 0; Mismatches 0; Indels
   Sequence 255, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
             Sequence 235, Application US/08467023
  1 KSMKVTVAFNQFG 13
   MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-467-023-235
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Gaps ö

Length 15; Indels

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COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
CORENATING SYSTEM: PC-DOS/MS-DOS
SOFTWARENT APRICATION DAIA:
APPLICATION DAIA:
APPLICATION NUMBER: US/08/467,023
FILING DAIT: June 6, 1995
CLASSIFICATION: 424
RIOR APPLICATION DAIA:
APPLICATION NUMBER: 08/350,225
FILING DAIT: December 6, 1994
ATTORNEY/AGENT INPORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMUNICATION NUMBER: 025.6 USD2 (IMI-028CPD2)
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 256:
SEQUENCE CHARACTERISTICS:
LEMOTH: 15 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: maino acide
TYPE: internal
MOLECULE TYPE: peptide
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US-08-467-023-256
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DD
2 KSMKYTVARNOPG 14
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 20
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  Scoring table:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

|           |        | Description     | Sequence 59, Appl | 9                | Sequence 61, Appl | . 28             |                     | Sequence 397. App | Seguence 16, Appl | 23,              | 16,              | 45               | 23               | 23               | Sequence 196, App |
|-----------|--------|-----------------|-------------------|------------------|-------------------|------------------|---------------------|-------------------|-------------------|------------------|------------------|------------------|------------------|------------------|-------------------|
| SUMMARIES |        | OI              | US-10-354-240-59  | US-10-354-240-60 | US-10-354-240-61  | US-10-354-240-58 | US-10-225-567A-1106 | US-09-834-765-397 | US-10-147-447-16  | US-10-300-072-23 | US-10-456-949-16 | US-10-456-947-45 | US-10-718-495-23 | US-10-717-984-23 | US-09-852-910-196 |
|           |        | DB              | 14                | 14               | 14                | 14               | 14                  | σ                 | 14                | 14               | 15               | 15               | 16               | 16               | 10                |
|           |        | Match Length DB | 15                | 15               | 15                | 15               | 19                  | 10                | 20                | 20               | 20               | 20               | 20               | 20               | 11                |
| *         | uery   | latch           | 72.5              | 67.9             | 48.6              | 46.8             | 32.1                | 30.3              | 30.3              | 30.3             | 30.3             | 30.3             | 30.3             | 30.3             | 28.4              |
|           | J.     | Score           | 79                | 74               | 53                | 51               | 32                  | 33                | 33                | 33               | 33               | 33               | 33               | 33               | 31                |
|           | Result | No.             | т                 | 7                | ო                 | 4                | ហ                   | 9                 | 7                 | 80               | 6                | 10               | 11               | 12               | 13                |

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Ouery Match 72.5%; Score 79; DB 14; Length 15; Best Local Similarity 93.3%; Pred. No. 1.1e-05; Matches 14; Conservative 0; Mismatches 1; Indels

1 QFGPNAGQRMPRARY 15

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PEATURE:
NAME/KEY: MISC PEATURE
COCATION: (1)...(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 45
US-10-354-240-59

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| RESULT 1 US-10-354-240-59 Sequence 59, A Septicator No APPLICANT: S APPLICANT: R APPLI | -240-9<br>e 59,<br>tion h<br>tion h<br>ANT:<br>ANT:<br>ANT:<br>ANT:<br>T FILLI<br>PILLING<br>OF IN<br>PILLING<br>OF SP | SULT 1  -10-354-240-59 Sequence 59, Application US/1035424 Publication No. US20030185847A1 APPLICANT: No. US20030185847A1 APPLICANT: Kume, Akinori APPLICANT: Kume, Akinori APPLICANT: Kume, Akinori APPLICANT: Iwama, Akiko APPLICANT: NUNEN: 2003-01-29 FILE REFERENCE: SPO-103D1 CURRENT APPLICATION NUMBER: US 09/14; PRIOR FILLING DATE: 1997-03-10 PRIOR FILLING DATE: 1997-03-10 PRIOR FILLING DATE: 1998-09-09 NUMBER OF SEQ ID NOS: 174 SEQ ID NO 59 LENGTH: 15 TWORE: PatentIn version 3:1 TWORE: DEPT | tion US 0301858 0301858 08hio khnori khnori Akiko Ohenke 10301 110301 110301 11997-03 11998-09-13 11998-09-13 | /103:<br>47A1<br>1 US<br>1 US<br>1 OT<br>1 OT<br>1 OS<br>0 OS<br>0 OS | ALIGNMENTS  10-354-240-59 equence 59, Application US/10354240 ublication No. US2030185847A1 ENERAL INFORMATION: APPLICANT: Sone, Toshio APPLICANT: Kume, Akinori APPLICANT: Iwama, Akinori APPLICANT: Wohsuke TITILE OF INVENTION: Peptide-Based Immunotherapeutic Agent FILE REFERENCE: SPO-103D1 CURRENT FILING DATE: 1997-003-10 PRIOR APPLICATION NUMBER: US 09/142,524 PRIOR APPLICATION NUMBER: US 09/142,524 PRIOR FILING DATE: 1998-09-09 PRIOR PRIOR PRIOR DATE: 1998-09-09 PRIOR PRIOR PRIOR DATE: 1998-09-09 | gent for Treating Allergic Dise.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |

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Sequence	106, Application US/10225567A
Sequence	106, Application US/10225567A
Sequence	106, Application US/20030113798A1
Sequence	106, Application No. US20030113798A1
GENERAL INFORMATION:	
GENERAL INFORMATION:	
APPLICANT: Brown, Joseph P.	
APPLICANT: Brown, Joseph P.	
APPLICANT: Brown, Christine L.	
APPLICANT: Brown, Christine L.	
TILE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS,	
TILE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS,	
CURRENT FILING DATE: 2000-12-19	
PRIOR FILING DATE: 2000-12-19	
NUMBER OF SEQ ID NOS: 2292	
SEQ ID NO 1106	
LENGTH: 19	
LENGTH: 19	
   APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103DI
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT PILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 58
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   Score 35; DB 14; Length 19;
Pred. No. 1.7e+02;
4; Mismatches 1; Indels
  Query Match
Best Local Similarity 90.0%; Pred. No. 0.36;
Matches 9; Conservative 0; Mismatches 1; Indels
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  , LOCATION: (1)..(15); OTHER INFORMATION: Cryjl peptide, Figure 1, Row 44 US-10-354-240-58
                         ö
Pred. No. 0.17;
1; Mismatches
  Sequence 58, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
  TYPE: PRT ORGANISM: Cryptomeria japonica
   32.1%;
50.0%;
     90.06;
   Query Match 32.1
Best Local Similarity 50.0
Matches 5; Conservative
     Best Local Similarity 90.0
Matches 9; Conservative
  1 OFGPNAGORM 10
   6 OFGPNCGORM 15
   11 PRARYGLIHV 20
  , ORGANISM: Homo
US-10-225-567A-1106
  US-10-225-567A-1106
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  JAPPLICANT: Sone, Toshio

APPLICANT: Sone, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Iwama, Akiko

APPLICANT: Iwama, Akiko

APPLICANT: Iwama, Akiko

APPLICANT: Iwama, Akiko

APPLICANT: Kino, Kohsuke

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: POTO-10301

CURRENT FILING DATE: 2003-01-29

PRIOR FILING DATE: 1997-03-10

PRIOR PILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFUTANTE: Patentin version 3.1

LENGTH: 15
  # PUDILICATION NO. USZUUJULBD84/AL

GENERAL INFORMATION:
# APPLICANT: Sone, Toshio
# APPLICANT: Kume, Akinori
# APPLICANT: Kume, Akinori
# APPLICANT: Wama, Akiko
# APPLICANT: Wama, Akiko
# APPLICANT: Wama, Akiko
# TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
# TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
# TITLE OF INVENTION: WUBBR: US/10/324,240
# CURRENT APPLICATION NUMBER: DCT/JP97/00740
# PRIOR FILING DATE: 1999-09
# PRIOR FILING DATE: 1999-09
# PRIOR FILING DATE: 1998-09-09
# NUMBER OF SEQ ID NOS: 174
# SOFTWARE: PatentIn version 3.1
# SEQ ID NO 60
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  DB 14; Length 15;
   Length 15;
   0, Indels
  LOCATION: (1).7(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 46
   cuery Match 67.9%; Score 74; DB 14;
Best Local Similarity 92.9%; Pred. No. 7.3e-05;
Matches 13; Conservative 1; Mismarches A
  LOCATION: (1)..(15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row
   48.6%; Score 53;
   US-10-354-240-61

'Sequence 61, Application US/10354240

'Publication No. US20030185847A1

GENERAL INFORMATION:
  US-10-354-240-60
; Sequence 60, Application US/10354240
; Publication No. US20030185847A1
  TYPE: PRT
ORGANISM: Cryptomeria japonica
  ORGANISM: Cryptomeria japonica
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   2 GORMPRARYGLVHV 15
   7 GORMPRARYGLIHV 20
  NAME/KEY: MISC_FEATURE
   NAME/KEY: MISC_FEATURE
  US-10-354-240-61
   US-10-354-240-60
   Query Match
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; Sequence 45, Application US/10456947; Publication No. US20040053841A1
  Query Match 30.3
Best Local Similarity 60.0
Matches 6; Conservative
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4 PNAPKRLPSA 13
  4 PNAGORMPRA 13
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   PNAPKRLPSA 13
   TYPE: PRT
ORGANISM: Homo sapiens
  TYPE: PRT
) ORGANISM: Homo Sapien
US-10-456-949-16
   US-10-456-949-16
  US-10-456-947-45
   LENGTH: 20
   Matches
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   Sequence 397, Application US/09834765
Fatent No. US2002005547841
Facent No. US2002005547841
Facent No. US2002005547841
Facent No. US2002005547841
Facent Nary Faria
Fapricant: Mary Faria
Fapricant: Arthur B. Raitano
Fapricant: Steve Chappell Mitchell
FAPLICANT: Aya Jakobovits
FILE REFERENCE: 129.6USU
FILE REFERENCE: 129.6USU
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
FRIOR FILING DATE: 2000-04-12
  Query Match 30.3%; Score 33; DB 14; Length 20; Best Local Similarity 60.0%; Pred. No. 3.8e+02; Matches 6; Conservative 2; Mismatches 2; Indels
   Length 10;
   4; Indels
   Sequence 16, Application US/10147447; Sequence 16, Application US/1014747; Publication No. US20330060410A1
GENERAL INFORMATION:
APPLICANT: Tracey, Kevin J.
APPLICANT: Yang, Huan
APPLICANT: Warren Jr., Howland Shaw
APPLICANT: Warren Jr., Howland Shaw
TITLE OF INVENTION: Use of HMG Fragments as
FITLE OF INVENTION: Anti-Inflammatory Agents;
FILE REFERENCE: 3268.1001-001
CURRENT APPLICATION NUMBER: US/10/147,447
CURRENT FILING DATE: 2002-08-16
PRIOR FILING DATE: 2001-05-15
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 20
   30.3%; Score 33; DB 9; I
60.0%; Pred. No. 1.9e+02;
tive 0; Mismatches 4;
  NUMBER OF SEQ ID NOS: 770
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 397
LENGTH: 10
   Query Match 30.3
Best Local Similarity 60.0
Matches 6; Conservative
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PNAPKRLPSA 13
  PNAGORMPRA 13
  3 GPNAGQRMPR 12
  1 GPPGGSRWPR 10
   4 PNAGQRMPRA 13
                                     8 PGSGQQLPRS 17
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US-09-834-765-397
  ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-147-447-16
   US-09-834-765-397
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US-10-300-072-23

Sequence 23, Application US/10300072

Publication No. US20030144201A1

APPLICANT: Howard Shaw Warren, Jr.

TITLE OF INVENTION: AGENTS

FILE OF INVENTION: AGENTS

FILE REFERENCE: 3269.1001-005

CURRENT APPLICATION NUMBER: US/10/300,072

CURRENT APPLICATION NUMBER: US 10/147,447

PRIOR PILING DATE: 2001-05-15

PRIOR FILING DATE: 2001-05-15

PRIOR APPLICATION NUMBER: US 60/291,034

PRIOR FILING DATE: 2001-05-15

NUMBER OF SEQ ID NOS: 58

SOFTWARE: FastSEQ for Windows Version 4.0

SED ID NO 23
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  Score 33; DB 14; Length 20;
Pred. No. 3.8e+02;
2; Mismatches 2; Indels
  Length 20;
   Ouery Match 30.3%; Score 33; DB 15; Length 20 Best Local Similarity 60.0%; Pred. No. 3.8e+02; Matches 5; Conservative 2; Mismatches 2; Indels
   Sequence 16, Application US/10456949

Publication No. US20040005316A1

GENERAL INFORMATION:

APPLICANT: Kevin J. Tracey

APPLICANT: Huan Yang

TITLE OF INVENTION: ANTI-INFLAMMATORY AGENTS

TITLE OF INVENTION: ANTI-INFLAMMATORY AGENTS

TITLE OF INVENTION OF A PROPER OF A PRIOR APPLICATION NUMBER: US/10/456,949

CURRENT FILING DATE: 2003-06-06

PRIOR APPLICATION NUMBER: 0/291,034

PRIOR PILING DATE: 2002-05-15

PRIOR FILING DATE: 2001-05-15

NUMBER OF SEQ ID NOS: 23

SOFTWARE: PRECED FOR WINDOWS VERSION 4.0

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TENGEN OF SEQ ID NOS: 23
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; ORGANISM: Homo sapiens
US-10-717-984-23
   US-10-411-336A-196
   RESULT 13
US-09-852-910-196
                      SEQ ID NO 23
LENGTH: 20
   RESULT 14
  Db
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                    GENERAL INFORMATION:

APPLICANT: Kevin J. Tracey
APPLICANT: Huan Yang
TITLE OF INVENTION: HHIBITORS OF THE INTERACTION BETWEEN
TITLE OF INVENTION: HWGB POLYPEPTIDES AND TOLL-LIKE RECEPTOR 2 AS
TITLE OF INVENTION: ANTI-INFLAMMATORY AGENTS
TITLE OF INVENTION: ANTI-INFLAMMATORY AGENTS
FILE REFERENCE: 3268 1001-007
CURRENT FILING DATE: 2003-06-06
PRIOR APPLICATION NUMBER: 10/147,447
PRIOR PILING DATE: 2002-05-15
PRIOR APPLICATION NUMBER: 60/291,034
PRIOR PILING DATE: 2001-05-15
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Pred. No. 3.8e+02;
2; Mismatches 2; Indels
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   Sequence 23, Application US/10717984
| Publication No. US20040156851A1
| GENERAL INFORMATION: Walter
| APPLICANT: Newman, Walter
| TITLE OF INVENTION: HMGBL COMBINATION THERAPIES
| FILE REFERENCE: 3258.1008-001
| CURRENT APPLICATION NUMBER: US/10/717,984
| CURRENT FILING DATE: 2003-11-20
| PRIOR FILING DATE: 2003-11-20
| PRIOR FILING DATE: 2003-11-20
| NUMBER OF SEQ ID NOS: 58
  Sequence 23, Application US/10718495
Sequence 23, Application US/10718495
Publication No. US20040141948A1
GENERAL INFORMATION:
APPLICANT: O'Keefe, Theresa L.
TITLE OF INVENTION: USE OF HWGB FRAGMENTS AS
TITLE OF INVENTION: ANTI-INFLAMMATORY AGENTS
FILE REFERENCE: 3258.1009-001
CURRENT APPLICATION NUMBER: US/10/718,495
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PRIOR APPLICATION NUMBER: 60/427,841
PRIOR PILING DATE: 2002-11-20
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Best Local Similarity 60.0
Matches 6; Conservative
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Best Local Similarity 60.0
Matches 6; Conservative
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ORGANISM: Homo sapiens
  TYPE: PRT
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US-10-717-984-23
   US-10-718-495-23
  US-10-456-947-45
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LENGTH: 20
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APPLICANT: Hammit dilchrist, Annette
APPLICANT: Hammit dilchrist, Annette
TITLE OF INVENTION: Method For Identifying Inhibitors of G Protein Coupled Receptor S
FILE REFERENCE: 2661-101
CURRENT FILIO DATE: 2001-09-18
PRIOR APPLICATION NUMBER: US 60/275,472
PRIOR APPLICATION NUMBER: US 60/275,472
SPRIOR FILING DATE: 2001-03-14
NUMBER OF SEQ ID NOS: 271
SOFTWARE: Patentin version 3.0
SEQ ID NO 196
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   28.4%; Score 31; DB 10; Length 11; 54.5%; Pred. No. 4.3e+02; tive 2; Mismatches 3; Indels
Score 33; DB 16; Length 20;
Pred. No. 3.8e+02;
2; Mismatches 2; Indels
  ; FEATURE:
; NAMEYER:
; LOCATION: (1)...(11)
; OTHER INFORMATION: Gs library peptide
US-09-852-910-196
  Sequence 196, Application US/09852910 Publication No. US20030096297A1 GENERAL INFORMATION:
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Best Local Similarity 60.0%;
Matches 6; Conservative 2
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  Query Match 28.4
Best Local Similarity 54.5
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   1 ORMPLROYELL 11
   4 PNAGQRMPRA 13
  ||| :|:| |
4 PNAPKRLPSA 13
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US-10-354-240-9

US-10-354-240-9

US-10-354-240-9

Sequence 9, Application US/10354240

Publication No. US20030185847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Rume, Akinori

APPLICANT: Rume, Akino

APPLICANT: Rume, Akinori

CURRENT Films OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION NUMBER: US/10/354,240

CURRENT FILING DATE: 1997-023-10

PRIOR APPLICATION NUMBER: US 09/142,524

PRIOR FILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFTWARE: PatentIn version 3.1

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; OTHER INFORMATION: Gs library peptide US-10-411-336A-196
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; ORGANISM: Cryptomeria japonica
US-10-354-240-9
  Query Match
Best Local Similarity 100.0
Matches 5; Conservative
  8 ORMPRARYGLI 18
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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1 PRARYGLIHVANNNYDPWSI 20
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

|           |            | tion        | nce 62, Appl     | nce 61, Appl     | nce 63, Appl     | nce 60, Appl     | nce 515, App      | nce 515, App      | nce 69, Appl      | nce 7, Appli    | nce 25, Appl     | nce 1274, Ap       | ce 34455, A         | nce 89, Appl     | Sequence 89, Appl |
|-----------|------------|-------------|------------------|------------------|------------------|------------------|-------------------|-------------------|-------------------|-----------------|------------------|--------------------|---------------------|------------------|-------------------|
|           |            | Description | Sequence         | Sequence         | Sequence         | Sequence         | Sequence          | Sequence          | Sequence          | Sequence        | Seguence         | Seque              | Seguen              | Seque            | Seque             |
| SUMMAKIES |            | OI          | US-10-354-240-62 | US-10-354-240-61 | US-10-354-240-63 | US-10-354-240-60 | US-10-012-542-515 | US-10-115-123-515 | US-10-300-694A-69 | US-10-468-543-7 | US-10-468-543-25 | US-10-813-638-1274 | US-09-864-761-34455 | US-10-269-695-89 | US-10-410-998-89  |
|           |            | ОВ          | 14               | 14               | 14               | 14               | 14                | 14                | 14                | 15              | 15               | 17                 | σ                   | 14               | 14                |
|           | *<br>Ouerv | Length      | 15               | 15               | 15               | 15               | 16                | 16                | 13                | 11              | 11               | 14                 | 19                  | 20               | 20                |
|           | Ouerv      | Match       | 72.2             | 71.3             | 50.4             | 46.1             | 32.2              | 32.2              | 30.4              | 28.7            | 28.7             | 28.7               | 28.7                | 28.7             | 28.7              |
|           |            | Score       | 83               | 82               | 58               | 53               | 37                | 37                | 35                | 33              | 33               | 33                 | 33                  | 33               | 33                |
|           | Result     | No.         | -                | 8                | m                | 4                | 'n                | 9                 | 7                 | 80              | đ                | 10                 | 11                  | 12               | 13                |

| ທີ່ສິດ                                             | Sequence 8, Appli<br>Sequence 12, Appl<br>Sequence 44, Appl                          | 1001                                                           | 29,<br>16,                                                      | Sequence 4s, Appli<br>Sequence 4, Appli<br>Sequence 4, Appli | Sequence or, Appl<br>Sequence 112, Appl<br>Sequence 81, Appl<br>Sequence 112, Appl     | w 9 1 -                   | 31, 7                                  | Sequence 13, Appl<br>Sequence 32, Appl<br>Sequence 49, Appl<br>Sequence 32, Appl<br>Sequence 76, Appl |
|----------------------------------------------------|--------------------------------------------------------------------------------------|----------------------------------------------------------------|-----------------------------------------------------------------|--------------------------------------------------------------|----------------------------------------------------------------------------------------|---------------------------|----------------------------------------|-------------------------------------------------------------------------------------------------------|
| US-10-100-608B<br>US-10-100-608B<br>US-10-099-791E | 6 US-10-099-791E-8<br>7 US-10-474-213-12<br>6 US-10-652-407-44<br>3 US-10-114-176-23 | 4 US-10-323-013-23<br>3 US-10-114-176-29<br>4 US-10-169-351-14 | 4 US-10-323-013-29<br>5 US-10-447-257-16<br>0 US-09-966-782A-48 |                                                              | 4 US-10-269-695-81<br>4 US-10-269-695-112<br>4 US-10-410-998-81<br>4 US-10-410-998-112 | 0000                      | 0-114-176-<br>0-032-221B<br>0-323-013- | ভ ক ক ক<br>⊃                                                                                          |
| 111                                                | 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7            |                                                                | 9042                                                            |                                                              | 2000                                                                                   | 7788                      | പ്പ്പ്<br>-തെത                         | 21112                                                                                                 |
| 27.                                                | 17 32 27.8<br>18 32 27.8<br>19 31 27.0<br>20 30 26.1                                 |                                                                | 0000                                                            | 28 30 26.1<br>29 30 26.1<br>30 30.26.1                       | 26.1<br>26.1<br>26.1<br>26.1                                                           | 25.7<br>25.7<br>25.2<br>2 | 1000                                   | 41 29 25.2<br>42 29 25.2<br>43 29 25.2<br>44 29 25.2<br>45 29 25.2                                    |

## ALIGNMENTS

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Sequence 62, Application US/10354240

Sequence 62, Application WJ/10354240

Publication No. US20030188847A1

GENERAL INFORMATION:

APPLICANT: Sone, Toshio

APPLICANT: Sone, Toshio

APPLICANT: Dairiki, Kazuo

APPLICANT: Iwama, Akiko

APPLICANT: Iwama, Akiko

FILE REFERENCE: SPO-103D1

CURRENT APPLICATION NUMBER: US/10/354,240

CURRENT FILING DATE: 1997-03-10

PRIOR FILING DATE: 1997-03-10

PRIOR PILING DATE: 1997-03-10

PRIOR FILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFTWARE: Patentin version 3.1

FEQUENCE: Patentin version 3.1

FEQUENCE: Patentin version 3.1
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   ; LOCATION: (1). (15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 48
US-10-354-240-62
  TYPE: PRT ORGANISM: Cryptomeria japonica
   NAME/KEY: MISC_FEATURE
   LENGTH: 15
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6 GLIHVANNNYDPWSI 20

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Gaps

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Sequence 60, Application US/10354240
; Sequence 60, Application US/10354240
; Publication No. US20030185847A1
; GENERAL INFORMATION:
    APPLICANT: Sone, Toshio
    APPLICANT: Kume, Akinori
    APPLICANT: Iwama, Akinori
    FILE REFERENCE: 2003-01-29
    CURRENT APPLICATION NUMBER: US/10/340
    PRIOR APPLICATION NUMBER: US/09/142,524
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    PRIOR PILING DATE: 1997-03-10
    PRIOR PILING DATE: 1998-09-09
    NUMBER OF SEQ ID NOS: 174
    SOFTWARE: PatentIn version 3.1
    IRANGTH: 18
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   60/089,510
                           ö
  Length 15;
  Indels
  US-U012-242-515

Sequence 515, Application US/10012542

Publication No. US20030044851A1

SEGUERAL INPORMATION:

APPLICANT: RUben et al.

TITLE OF INVENTION: 94 Human Secreted Proteins

FILE REFERENCE: P2029P1

CURRENT FILING DATE: 2001-12-12

CURRENT FILING DATE: EARLIER APPLICATION NUMBER: 09/49

PRIOR FILING DATE: EARLIER APPLICATION NUMBER: 60/69

PRIOR PILING DATE: EARLIER FILING DATE: 1999-12-14

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/69

PRIOR PILING DATE: EARLIER FILING DATE: 1998-06-16

PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-16

PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-16

PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-16

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/6

PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1998-06-22

PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1998-06-22

PRIOR PRIOR EARLIER FILING DATE: 1998-06-22

NUMBER OF SEQ ID NOS: 532

SOFTWARE: PATEURI VET. 2.0
                           Indels
  , LUCATION: (1). T(15)
, OTHER INFORMATION: Cryjl peptide, Figure 1, Row 46
US-10-354-240-60
   ö
                              ;
   46.1%; Score 53; DB 14; 90.0%; Pred. No. 0.43;
90.0%; Pred. No. 0.075;
tive 1; Mismatches
  1; Mismatches
   ORGANISM: Cryptomeria japonica
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Section 9, Conservative
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   1 PRARYGLIHV 10
  1 ANNNYDPWTI 10
  11 ANNNYDPWSI 20
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Matches 9; Conserv
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  g
  GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Kume, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Kume, Akinori

APPLICANT: Twama, Akiko

APPLICANT: Twama, Akiko

APPLICANT: Twama, Akiko

APPLICANT: Twama, Rohsuke

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea

TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Disea

CURRENT APPLICANTION NUMBER: US/10/354,240

PRIOR FILING DATE: 1997-03-10

PRIOR FILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFTWARE: Patentin version 3.1

LENGTH: 15
   TITLE OF INTENTAL ALLOW. Peptide-Based Immunotherapeutic Agent for Treating Allergic Diseating FILE REFERENCE: SPO-103D1

CURRENT APPLICATION NUMBER: US/10/354,240

CURRENT PILING DATE: 1030-301-29

PRIOR APPLICATION NUMBER: US 09/142,524

PRIOR FILING DATE: 1997-03-10

RIOR APPLICATION NUMBER: US 09/142,524

PRIOR FILING DATE: 1998-09-09

NUMBER OF SEQ ID NOS: 174

SOFTWARE: Patentin version 3.1
   ô
  Gaps
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   DB 14; Length 15;
   Length 15;
   0; Indel's
   LOCATION: (1). (15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 47
   Query Match 71.3%; Score 82; DB 14; Best Local Similarity 93.3%; Pred. No. 1.7e-05; Matches 14; Conservative 1; Mismatches n.
   LOCATION: (1). (15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row
  50.4%; Score 58;
  Sequence 53, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
  Sequence 61, Application US/10354240
Publication No. US20030185847A1
   TYPE: PRT ORGANISM: Cryptomeria japonica
  ORGANISM: Cryptomeria japonica
   1 PRARYGLIHVANNNY 15
   1 GLVHVANNNYDPWTI 15
  APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
   APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
  NAME/KEY: MISC_FEATURE
   NAME/KEY: MISC_FEATURE
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   US-10-354-240-61
   SEQ ID NO 63
LENGTH: 15
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   32.2%; Score 37; DB 14; Length 16; 55.6%; Pred. No. 1.3e+02; tive 2; Mismatches 2; Indels
  Score 37; DB 14; Length 16;
Pred. No. 1.3e+02;
2; Mismatches 2; Indels
   2; Indels
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US-10-115-123-515

Squence 515, Application US/10115123

Publication No. US20030065151A1

GENERAL INFORMATION:

TITLE OF INVENTION: 94 Human Secreted Proteins
FILE REFERENCE: PZ029G30AP1D2

CURRENT APPLICATION NUMBER: US/10/115,123

CURRENT FILING DATE: 2002-04-04

PRIOR FILING DATE: 1999-06-15

PRIOR FILING DATE: 1999-06-15

PRIOR FILING DATE: 1999-06-16

PRIOR FILING DATE: 1998-06-16

PRIOR PRIOR PRIOR DATE: 1998-06-16

PRIOR PRIOR DATE: 1998-06-16

   Sequence 69, Application US/10300694A
Publication No. US20030185870A1
GENERAL INFORMATION:
APPLICANT: UNFORMATION:
APPLICANT: Grinstaff, Mark W.
APPLICANT: Malah, Daniel J.
APPLICANT: Walsh, Elisabeth B.
APPLICANT: Middleton, Crystan
ITLE OF INVENTION: INVERFACIAL BIOMATERIALS
FILE REFERENCE: 180/143/2
CURRENT FILING DATE: 2003-05-07
  CURRENT FILING DATE: 2003-05-07
PRIOR PRING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 117
  Query Match
Best Local Similarity 55.6%;
Matches 5; Conservative
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Best Local Similarity 55.0
-hea 5; Conservative
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CORGANISM: Homo sapiens
US-10-012-542-515
   : | | | | | | 5
5 SNGNLDPWA 13
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US-10-115-123-515
   11 ANNNYDPWS 19
   US-10-300-694A-69
SEQ ID NO 515
LENGTH: 16
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Sequence 25, Application US/10468543

Sequence 25, Application US/10468543

Sequence 25, Application No. US20040091938A1

GENERAL INFORMATION:
APPLICANT: Infunxa, Tatsuro
APPLICANT: Win, Mijung
APPLICANT: Win, Mijung
APPLICANT: Ono, Takashi
ITILE OF INVENTION: Lectins for Analyzing Sugar Chains and Method of Using the Same
FILE REFERENCE: 03-786
CURRENT APPLICATION NUMBER: US/10/468,543
CURRENT APPLICATION NUMBER: US 2003-08-20
PRIOR PILING DATE: 2001-02-20
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn version 3.1
SEQ ID NO 25
LENGTH: 11
  APPLICANT: Itimura, Tatsuro
APPLICANT: Itimura, Tatsuro
APPLICANT: Misumoto, Mariko
APPLICANT: Yim, Mijung
APPLICANT: Takashi
TITLE OF INVENTION: Lectins for Analyzing Sugar Chains and Method of Using the Same FILE REFERENCE: 03-786
CURRENT APPLICATION WUMBER: US/10/468,543
CURRENT FILING DATE: 2003-08-20
PRIOR FILING DATE: 2001-02-20
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  Score 33; DB 15; Length 11;
Pred. No. 3.5e+02;
2; Mismatches 0; Indels
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   ; OTHER INFORMATION: Polycarbonate-binding peptide 69
US-10-300-694A-69
  Query Match 30.4%; Score 35; DB 14; L
Best Local Similarity 71.4%; Pred. No. 2.1e+02;
Matches 5; Conservative 1; Mismatches 1;
  US-10-468-543-7

; Sequence 7, Application US/10468543

; Publication No. US20040091938A1

; GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 69
LENGTH: 13
   NUMBER OF SEQ ID NOS: 95
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SEQ ID NO 7
  LENGTH: 11
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US-10-468-543-7
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US-10-468-543-25
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Best Local Similarity 66.7
Matches 4; Conservative
  12 NNNYDPW 18
  TYPE: PRT ORGANISM: Artificial
  3 NVNYNPW 9
  13 NNYDPW 18
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6 HSYDPW 11
   FEATURE:
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US-10-269-695-89

Sequence 89, Application US/10269695

Publication No. 1220030229023A1

Publication No. 1220030229023A1

APPLICANT: OLINER, JONATHAN DANIEL

APPLICANT: OLINER, JONATHAN DANIEL

APPLICANT: OLINER, JONATHAN DANIEL

FILE REFERENCE: A-801A

TITLE OF INVENTION: SPECIFIC BINDING AGENTS OF HUMAN ANGIOPOIETIN-2

FILE REFERENCE: A-801A

TITLE OF INVENTION SPECIFIC BINDING AGENTS OF HUMAN ANGIOPOIETIN-2

FILE REFERENCE: A-801A

FILE REFERENCE: A-801A

FILE REPERENCE: A-801A

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   TYPE: PRT ACCEPANT ARTIFICIAL Sequence PEATURE: OTHER INFORMATION: Polypeptide capable of binding to Ang-2
  OTHER INFORMATION: EXPRESSED IN HELLOO, SIGNAL = 3.8

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 5

OTHER INFORMATION: EXPRESSED IN BEALN, SIGNAL = 4.3

OTHER INFORMATION: EXPRESSED IN BEARY, SIGNAL = 3.6

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 3.7

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Pred. No. 6.1e+02;
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00670
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 99/608,408
PRIOR PILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-06-30
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PRIOR PILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
   4; Mismatches
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APPLICATION NUMBER: PCT/US01/00665
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3 KRNYGILHIA 12
   2 RARYGLIHVA 11
   TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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Best Local Similarity
Matches 4; Conserv
  OTHER INFORMATION: NOTHER INFORMATION: FOTHER INFORMATION: COTHER INFORMATION: FOTHER 
  ; OTHER INFORMATIOUS-09-864-761-34455
   SEQ ID NO 34455
LENGTH: 19
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   g
  GENERAL INCRMATE Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENITON: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENITON: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aconica-X-1
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-26
PRIOR FILING DATE: 2000-06-36
PRIOR PILING DATE: 2000-06-36
PRIOR FILING DATE: 2000-06-36
PRIOR FILING DATE: 2000-06-36
PRIOR FILING DATE: 2000-09-37
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
   FIGUREAL INFORMATION:
APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin D.
TITLE OF INVENTION: USE THEREOF
FILE OF INVENTION: USE THEREOF
FILE REPERENCE: 15966-599
CURRENT APPLICATION NUMBER: US/10/813,638
CURRENT APPLICATION NUMBER: G0/163,783
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1468
SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 1274
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Score 33; DB 15; Length 11.
Pred, No. 3.5e+02;
  Sequence 34455, Application US/09864761 Patent No. US20020048763A1
   ; Sequence 1274, Application US/10813638; Publication No. US20040235026A1
  NAME/KEY: VARIANT
1 LOCATION: (7)...(0)
2 OTHER INFORMATION: CSNP translation
US-10-813-638-1274
  Query Match 28.7
Best Local Similarity 66.7
Matches 4; Conservative
  ORGANISM: Homo sapiens
  14 NYDPWSI 20
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6 SYNPWSL 12
  13 NNYDPW 18
  ::||||
6 HSYDPW 11
   GENERAL INFORMATION
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US-09-864-761-34455
   -10-813-638-1274
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6 SNNNFLTWS 14

US-10-269-695-89

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| Sequence 8, Application US/10100608B | Publication No. US20030104412A1 |
| GENERAL INFORMATION: | APPLICANT: Heiskala, Marja |
| TILE OF INVENTION: REG-LIKE PROTEIN |
| FILE REFERENCE: CDS-261 |
| CURRENT APPLICATION UNMBER: US/10/100,608B |
| CURRENT FILING DATE: 2002-09-10 |
| PRIOR APPLICATION NUMBER: 60/276,414 |
| PRIOR FILING DATE: 2002-03-16 |
| NUMBER OF SEQ ID NOS: 45 |
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   6 SNNNFLTWS 14
  TYPE: PRT
ORGANISM: Human
                          RESULT 15
US-10-100-608B-8
  US-10-100-608B-8
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   NS-10-410-998-89

Sequence 89, Application US/10410998

Sequence 89, Application US/10410998

Publication No. 1020030236193A1

GENERAL INFORMATION:

APPLICANT: OLINER, JONATHAN DANIEL

APPLICANT: OLINER, JONATHAN DANIEL

APPLICANT: MIN, HOSUNG

TITLE REFERENCE: A-801A

CURRENT PAPLICATION NUMBER: US/10/410,998

CURRENT PILING DATE: 2003-04-09

PRIOR APPLICATION NUMBER: US/0/69,695

PRIOR PILING DATE: 2002-10-10

PRIOR PILING DATE: 2002-10-10

PRIOR PILING DATE: 2000-09-27

PRIOR PILING DATE: 2001-10-11

NUMBER OF SEQ ID NOS: 359

SOFTWARE: PARCHIN VERSION 3.1

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2; Mismatches 0; Indels
  Length 20;
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28.7%; Score 33; DB 14;
Best Local Similarity 66.7%; Pred. No. 6.4e+02;
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Sequence 5, Application US/10100608B
PUDLICATION NO. US20030104412A1
GENERAL INFORMATION:
TITLE OF INVENTION: REG-LIKE PROTEIN
FILLE REFERENCE: CDS-261
CURRENT PELICATION NUMBER: US/10/100,608B
CURRENT FILING DATE: 2002-09-10
FRIOR APPLICATION NUMBER: 60/276,414
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FRIOR APPLICATION NUMBER: 50/276,414
SRIOR FILING DATE: 2002-03-16
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Best Local Similarity 66.7
Matches 4; Conservative
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8 NWDPWT 13
  14 NYDPWS 19
  14 NYDPWS 19
  TYPE: PRT
ORGANISM: Human
  US-10-100-608B-5
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11 ANNNYDPWS 19

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27.8%; Score 32; DB 14; Length 17; 55.6%; Pred. No. 7.7e+02; tive 2; Mismatches 2; Indels

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LENGTH: 20 amino acids
TYPE: amino acid
TOPOLOGY: linear
  MOLECULE TYPE: peptide FRAGMENT TYPE: internal
 88668800661111111
USA
   STATE: MA
COUNTRY: U
   ZIP: 02154
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US-08-467-023-49
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   January 26, 2005, 15:54:02; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
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(c) 1993 - 2005 Compugen Ltd.
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  SUMMARIES
  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                              - protein search, using sw model
   1 PRARYGLIHVANNNYDPWSI 20
  BLOSUM62
Gapop 10.0 , Gapext 0.5
  US-09-202-464-26
  Query
Match Length
           Copyright
   Minimum DB seq length: 0 Maximum DB seq length: 20
  <u>ორ</u> 4 ო ტ
  Title:
Perfect score:
  Scoring table:
                              OM protein
  Database :
   Sequence:
   Searched:
   Run on
  Result
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US-09-142-524D-61
US-09-142-524D-62
  LENGTH: 15
   g
   ð
  ö
   ö
  Gaps
   Gaps
  ö
   ö
  APPLICANT: Griffeth, Irwin J.;
APPLICANT: Bollock, Joanne;
APPLICANT: Bollock, Joanne;
APPLICANT: Bollock, Joanne;
APPLICANT: Garman, Richard D;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Exlay, Mark A.;
APPLICANT: Brauer, Andrew;
APPLICANT: Braver, Andrew;
APPLICANT: Convers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Appanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
  Score 104; DB 3; Length 19;
Pred. No. 1.6e-09;
2; Mismatches 0; Indels
   96.5%; Score 111; DB 3; Length 20; 90.0%; Pred. No. 1.4e-10; tive 2; Mismatches 0; Indels
   COMPUTE: MA

CONTUTER: MA

ZIP: 02154

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Date of the compatible
APPLICATION NUMBER: US/08/467,023
FILING DATE: Date of the compatible
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGRYT INPORMATION:
NAME: Jane E. Remillard
REFERENCE/DOCKET NUMBER: 38,872
REFERENCE/DOCKE
   Sequence 258, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
  1 RARYGLVHVANNNYDPWTI 19
   1 PRARYGLIHVANNNYDPWSI 20
  1 PRARYGLVHVANNNYDPWTI 20
   2 RARYGLIHVANNNYDPWSI 20
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Best Local Similarity 89.5%;
Matches 17; Conservative ;
   Query Match
Best Local Similarity 90.09
Matches 18, Conservative
  MOLECULE TYPE: peptide FRAGMENT TYPE: internal
  amino acid
  US-08-467-023-258
  RESULT 2
US-08-467-023-258
            US-08-467-023-49
  ò
   ò
   g
```

RESULT 3

셤

```
US-US-144-544D-54

j Sequence 62, Application US/09142524D

patent No. 6719976

j GENERAL INFORMATION:
    APPLICANT: Sone, Toshio
    APPLICANT: Wime, Akinori
    APPLICANT: 1998-09-09
    PRIOR APPLICATION NUMBER: PCT/JP97/00740
    PRIOR APPLICATION NUMBER: PCT/JP97/00740
    NUMBER OF SEQ ID NOS: 174
    SOFTWARE: PatentIn version 3.1
    SEQ ID NO 62
    LENGTH: 15
    TYPE: PRT
    TYPE: PRT
    CORGANTSM: CTyptomeria japonica
  GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: State, Arinori
APPLICANT: Mame, Arinori
APPLICANT: Iwama, Ariko
APPLICANT: Iwama, Ariko
APPLICANT: Iwama, Ariko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disease
TITLE REPRENCE: SPO-103
CURRENT FILING DATE: 1988-09-09
PRIOR FILING DATE: 1988-09-09
PRIOR FILING DATE: 1989-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
FERRALL OF THE SEC ID NOS: 174
SOFTWARE: Patentin version 3.1
  ö
  ö
  Gaps
  Gaps
  ö
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  Length 15;
  72.2%; Score 83; DB 4; Length 15; 86.7%; Pred. No. 2.1e-06; tive 2; Mismatches 0; Indels
  Indels
   ; LOCATION: (1).7(15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 47
US-09-142-524D-61
   Score 82; DB 4; Le:
Pred. No. 2.9e-06;
1; Mismatches 0;
  FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (1)...(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 48
US-09-142-524D-62
   ; Sequence 61, Application US/09142524D
; Patent No. 6719976
  TYPE: PRT
ORGANISM: Cryptomeria japonica
   71.3%;
93.3%;
   1 PRARYGLVHVANNNY 15
   1 PRARYGLIHVANNNY 15
   6 GLIHVANNNYDPWSI 20
   1 GLVHVANNNYDPWTI 15
   Query Match 71.3
Best Local Similarity 93.3
Matches 14; Conservative
   FEATURE:
NAME/KEY: MISC_FEATURE
  g
   ò
```

```
APPLICANT: Some, Toshio
APPLICANT: Some, Toshio
APPLICANT: Some, Akinori
APPLICANT: Name, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 60
   ö
   ö
  Gaps
   Gaps
   ö
   ö
  Score 58; DB 3; Length 20;
Pred. No. 0.02;
1; Mismatches 0; Indels
   Length 15;
  0; Indels
  025.6 USD2 (IMI-028CPD2)
   ; NAME/KEY: MISC_FEATURE
; LOCATION: (1)...(15)
... OTHER INFORMATION: Cryjl peptide, Figure 1, Row 46
US-09-142-524D-60
  Query Match 46.1%; Score 53; DB 4;
Best Local Similarity 90.0%; Pred. No. 0.086;
Matches 9; Conservative 1; Mismatches
   US-09-142-524D-60
; Sequence 60, Application US/09142524D
; Patent No. 6719976
  ; Sequence 48, Application US/08467023
; Patent No. 6090386
  GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, mai-Chang;
APPLICANT: Yeung, Siu-mei H.;
                               NAME: Jane E. Kemillala
REGISTRATION UNDRER: 38,972
REPRENCE/DOCKET NUMBER: 025.6
TELECOMMUNICATION INFORMATION:
TELEFHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
  ORGANISM: Cryptomeria japonica
  Query Match 50.4%;
Best Local Similarity 90.0%;
Matches 9; Conservative 1
          ATTORNEY/AGENT INFORMATION:
   MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-467-023-50
  11 ANNNYDPWSI 20
   1 ANNNYDPWTI 10
  1 PRARYGLIHV 10
   amino acid
  linear
   Patent No. 6719976
GENERAL INFORMATION:
   TOPOLOGY:
   US-08-467-023-48
  LENGTH: 15
  TYPE: PRT
  ઠે
  d
  ઠે
  셤
   GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Lauriki, Kazuo
APPLICANT: Ivama, Akinori
APPLICANT: Ivama, Akiko
APPLICANT: Nino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
PRIOR FILING DATE: 1998-09-09
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: PatentIn version 3.1
SEQ ID NO 63
  .;
0
  Gaps
  sequence 50, Application US/08467023

Patent No. 609036

GENERAL INFORMATION:

APPLICANT: Griffeth, Irwin J.;

APPLICANT: Bond, Unlian F.;

APPLICANT: Garman, Richard D;

APPLICANT: Keung, Markang;

APPLICANT: Renug, All-mei H.;

APPLICANT: Brauer, Andrew;

APPLICANT: Bowers, Steven P.;

TITLE OF INVENTION: Allergenic Proteins And Peptides From NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: ImmuLogic Pharmacceutical Corporation, Inc.

STREET: Glo Lincoln St.

COUTTY: Waltham

STATE: MA

COUNTRY: USA

TID.
  ;
0
  50.4%; Score 58; DB 4; Length 15; 90.0%; Pred. No. 0.015;
  0; Indels
  FEATURE:

NAME/KEY: MISC_FEATURE

COCATION: (1) __(15)

OTHER INPORMATION: Cryjl peptide, Figure 1, Row 49
US-09-142-524D-63
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   1; Mismatches
  US/08/467,023
  Sequence 63, Application US/09142524D
Patent No. 6719976
  08/350,225
  APPLICATION NUMBER: US/08/467
FILING DATE: June 6, 1995
CLASIPRICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,22:
FILING DATE: December 6, 1994
   TYPE: PRT
ORGANISM: Cryptomeria japonica
   Best Local Similarity 90.0
Matches 9; Conservative
  1 ANNNYDPWTI 10
  11 ANNNYDPWSI 20
   ZIP: 02154
RESULT 5
US-09-142-524D-63
  US-08-467-023-50
  Query Match
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COUNTRY: USA

ZIP: 02109-1875

COUNTRY: USA

COMPUTER READBLE FORM:

MEDIUW TYPE: Floppy disk

COMPUTER: IBM FC compatible

OPERATING SYSTEM: FC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/290,448A

FILING DATE: August 15, 1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/529,951

FILING DATE: March 17, 1989

ATTORNEY/AGENT INFORMATION:

NAME: Amadragouras

REGISTRATION NUMBER: 36,207
   Mismatches
  NUMBER OF SEQUENCES: 93
CORRESPONDENCES ADDRESSS:
AADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, suite 510
  IMI-018CN
  REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET UNMBER: IM
TELECOMMUNICATION: TELEPHONE: (617)227-7400
   TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
  Query Match
Best Local Similarity 61.5%;
Matches 8; Conservative
   TELEFAX: (617)227-5941
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
   : 17 amino acids
amino acid
3Y: linear
  6 GLIHVANNNYDPW 18
  1 GFFQVVNNNYDRW 13
   MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-290-448A-55
  CITY: Boston
STATE: Massachusetts
  TOPOLOGY:
   RESULT 10
   à
   ö
   Sequence 55, Application US/08290448A

Sequence 55, Application US/08290448A

Patent No. 567654

GENERAL INFORMATION:
APPLICANT: Ragner, David G.
APPLICANT: Ragner, David G.
APPLICANT: Rafner, Thorunn
APPLICANT: Rafner, Suit-C 510
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, suite 510
CITY: Boston
STREET: 60 State Street, suite 510
CITY: Boston
STREET: Massachusetts
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PLOPS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
  46.1%; Score 53; DB 3; Length 20; 90.0%; Pred. No. 0.12; cive 1; Mismatches 0; Indels
  COMPUTEX: USA

ZIP: OCHETICS: USA

ZIP: OCHETICS: COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane B. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION STORE NUMBER: 38,872
REGISTRATION STORE NUMBER: 38,872
REGISTRATION STORE NUMBER: 38,872
REGISTRATION STORE NUMBER: 38,872
RELEPAX: (617) 227-7400
FILLEPAX: (617) 227-740
  Query Match
Best Local Similarity 90.0
Matches 9; Conservative
   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
  1 PRARYGLIHV 10
  |||||||||||
11 PRARYGLVHV 20
   TYPE: amino acid
TOPOLOGY: linear
   STATE: MA
  US-08-290-448A-55
   US-08-467-023-48
   COUNTRY:
  ò
```

```
Gaps
  Segmence 55, Application US/08290448A
Factor No. 5698204
BENERAL INFORMATION:
APPLICANT: Rapper, David G.
APPLICANT: Rlapper, David G.
APPLICANT: Rlapper, David G.
APPLICANT: Rapper, David G.
APPLICANT: Applicant Thorun
APPLICANT: And Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
   ;
0
  Score 50; DB 1; Length 17; Pred. No. 0.28;
   5; Indels
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: August 15, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: March 17, 1989
ATTORNEY,AGENT INPORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-018CN
TELECOMUNICATION INPORMATION:
TELECOMUNICATION INPORMATION:
TELECOMUNICATION INPORMATION:
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amino acid
  Query Match
Best Local Similarity
US-08-461-939B-55
  US-08-461-939B-55
  US-08-464-000-55
  Best Loc
Matches
  g
   à
  ö
  Gaps
   ö
   APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Rafnar, Thorunn
APPLICANT: Ruo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
  ö
  DB 1; Length 17;
   43.5%; Score 50; DB 1; Length 17; 61.5%; Pred. No. 0.28; tive 0; Mismatches 5; Indels
  ADDRESSER LAHIVE & COCKFIELD, LLP
STREET: 60 State Street
CITY: Boston
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: Maxch 17, 1989
ATTORNEY/AGENT INFORMATION:
NAMME: Amy E. Manch gouras
RESTERRATION NUMBER: IMI-018DV
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
   Mismatches
  Query Match 43.5%; Score 50; Best Local Similarity 61.5%; Pred. No. (Matches 8; Conservative 0; Mismatch
   US-08-175-069A-55; Sequence 55, Application US/08175069A; Patent No. 5776761; GENERAL INFORMATION:
   TELEPAX: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
  6 GLIHVANNNYDPW 18
17 amino acids
   1 GFFQVVNNNYDRW 13
   6 GLIHVANNNYDPW 18
  17 amino acids
   GFFQVVNNNYDRW 13
   Query Match
Best Local Similarity 61.5
Matches 8; Conservative
   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
  amino acid
   US-08-290-448A-55
  US-08-175-069A-55
                                  FOPOLOGY:
   TOPOLOGY:
  LENGTH:
  ઠે
   g
   ઠે
```

RESULT 12

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Sequence 55, Application US/08461939B
Patent No. 6335019
GENERAL INFORMATION:
APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Rufnar, Thorun
APPLICANT: Rufnar, Thorun
APPLICANT: Rufnar, Thorun
APPLICANT: Rufnar, Protein
TITLE OF INVENTION: Methods For Treating Sensitivity To A
TITLE OF INVENTION: Protein Allergen Using Peptides Which Inlcude A T Cell Epitop
NUMBER OF SEQUENCES: 93
CORRESPONDENCES: 1
ANDRESSES:
ADDRESSES:
ADDRESSES:
ATREET: 28 State Street
CITY: Boston
  Sequence 55, Application US/08464000
Patent No. 6335020
GENERAL INFORMATION:
APPLICANT: Klapper, David G.
APPLICANT: Rafnar, Thorunh
APPLICANT: Rafnar, Thorunh
APPLICANT: Ro, Mei-chang
TITLE OF INVENTION: Allergenic Peptides from Ragweed Pollen
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
   Score 50; DB 3; Length 17; Pred. No. 0.28;
  Indels
   SCENARIA STATEMENT SELECTION #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,939B
  FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/464,000
FILING DATE: 05-UN-1995
FILING DATE: 15-MQ-1994
APPLICATION NUMBER: US 07/29,951
FILING DATE: 29-MAY-1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: 17-MAR-1989
APPLICATION NUMBER: US 07/325,365
FILING DATE: 17-MAR-1980
  LAHIVE & COCKFIELD, LLP
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET UNBER: IM
TELECOMMUNICATION:
TELEPHONE: (617)227-7400
   43.5%;
61.5%;
  TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
   6 GLIHVANNNYDPW 18
   GFFQVVNNNYDRW 13
  Conservative
   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
   Massachusetts
  COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  ADDRESSEE:
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1;
   Sequence 28, Application US/08290448A

Sequence 28, Application US/08290448A

Patent No. 5698204

GENERAL INFORMATION:

APPLICANT: Radper, David G.

APPLICANT: Radper, David G.

APPLICANT: Rafnar, Thorunn

APPLICANT: Rafnar, Thorunn

TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses

CORRESPONDENCES: 93

NUMBER OF SEQUENCES:
   Score 40.5; DB 1; Length 15; Pred. No. 7.2;
  Indels
  ADDRESSENCE ASTACE; SUITE 510
CITY: BOSCON
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IB PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 07/829,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: MA-CH 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: AMD SMACH 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: AMD SMACH 10, 1980
REGISTRATION NUMBER: IMI-018CN
TELECOMMUNICATION INFORMATION:
   2; Mismatches
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: MARCH 17, 1989
ATTORNEY/GENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REPERDICE/DOCKET NUMBER: IMI-018CN
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-7401
TELEPHONE: (617)227-7401
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
   TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
   Query Match
Best Local Similarity 50.0%;
Matches 8; Conservative ;
  1 PRARYGLIHVANNNYD 16
  || |:| : || ||
1 PR-RFGFFQIVNNFYD 15
  TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
   TOPOLOGY: linear
MOLECULE TYPE: peptide
PRAGMENT TYPE: internal
  LENGTH: 15 amino TYPE: amino acid
  TELEPHONE:
   US-08-290-448A-28
  US-08-290-448A-28
  В
  ö
  WESULY 148A-28

Sequence 28, Application US/08290448A

Patent No. 5676954

GENERAL INFORMATION:

APPLICANT: Righer, David G.

APPLICANT: Rafnar, Thornan

STATE: ABSACHUBER: COCKFIELD

STATE: Massachusetts

CONTYT: BOSTON

CONTYT: BOSTON

SOFTWARE: Patentin Release #1.0, Version #1.25

COMPUTER: Patentin Release #1.0, Version #1.25

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: US/08/290,448A

FILNG DATE: AUGUST 15, 1994

PRIOR APPLICATION DATE: AUGUST 15, 1994
  DB 3; Length 17;
   5; Indels
   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,000
FILING APPLICATION DATA:
APPLICATION NUMBER: US 08/290,448
FILING APPLICATION NUMBER: US 07/529,951
FILING DATE: 15-AUG-1994
APPLICATION NUMBER: US 07/529,951
FILING DATE: 25-MAY-1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: 17-MAR-1999
ATTORNEY AGENT INFORMATION:
NAME: AMW E. Mandragouras
REGISTRATION NUMBER: 1M1-018CN2
FELEPHONE: (617)227-740
TELEPHONE: (617)227-5941
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TWENDER: Amino acids
  Score 50; DB 3
Pred. No. 0.28;
0; Mismatches
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
   Query Match
Best Local Similarity 61.5%;
Matches 8; Conservative
              60 State Street
   6 GLIHVANNNYDPW 18
  1 GFFQVVNNNYDRW 13
                                       Boston
: Massachusetts
RY: USA
   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-08-464-000-55
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US-08-290-448A-28

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Query Match
Best Local Similarity 50.0%; Pred. No. 7.2;
Matches 8; Conservative 2; Mismatches 5; Indels

Qy 1 PRARYGLIHVANNNYD 16
| | | | | | | | | |
| Db 1 PRARYGLIHVANNNYD 15
| | | | | | | |
| Db 1 DRARYGLIHVANNNYD 15
| | | | | | |
| Db 1 DRARYGREPQINNNYD 15
| | | | | | |
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1; Gaps

Lyis bade Blank (nsbjo)

us-09-202-464-27.closed.rapb

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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
  Published Applications AA:*

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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| 7      | 82    | 71.9       | 15              | 14 | US-10-354-240-64   | Sequence 64, Appl |
| m      | 28    | 50.9       | 15              | 14 | US-10-354-240-62   | Sequence 62, Appl |
| 4      | 53    | 46.5       | 15              | 14 | US-10-354-240-65   | Sequence 65, Appl |
| 2      | 38    | 33.3       | 16              | 14 | US-10-346-162-100  | Sequence 100, App |
| 9      | 37    | 32.5       | 16              | 14 | US-10-012-542-515  | Sequence 515, App |
| 7      | 37    | 32.5       | 16              | 14 | US-10-115-123-515  | 515,              |
| ۵      | 35    | 30.7       | 13              | 14 | US-10-300-694A-69  | Sequence 69, Appl |
| σ      | 35    | 30.7       | 15              | 14 | US-10-346-162-142  | Sequence 142, App |
| 10     | 33    | 28.9       | 11              | 15 | US-10-468-543-7    | Sequence 7, Appli |
| 11     | 33    | 28.9       | 11              | 15 | US-10-468-543-25   | Sequence 25, Appl |
| 12     | 33    | 28.9       | 14              | 17 | US-10-813-638-1274 | Sequence 1274, Ap |
| 13     | 33    | 28.9       | 15              | 15 | US-10-182-936A-191 | Sequence 191, App |

| Sequence 275, App<br>Sequence 276, App | 8                | 89,    | 'n     | Sequence 8, Appli | 'n     | 8      | N      | 4                | m    | _      | П      | _         | œ        | æ         |           | σ      | N    | Ф      | 47,              | 23,        | Sequence 23, Appl | 29,    | 14, | 29,    | 16,      | 96     | 33,      | 141                | Sequence 2941, Ap | 294                |
|----------------------------------------|------------------|--------|--------|-------------------|--------|--------|--------|------------------|------|--------|--------|-----------|----------|-----------|-----------|--------|------|--------|------------------|------------|-------------------|--------|-----|--------|----------|--------|----------|--------------------|-------------------|--------------------|
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| 28                                     | 6.8              | 3 28.9 | 2 28.1 | 2 28.1            | 2 28.1 | 2 28.1 | 1 27.  | 1 27.2           | 27.2 | 1 27.2 | 1 27.2 | 1 27.2    | 1 27.2   | 7.2       | 5 26.     | 5 26.8 | 0 26 | 0 26   | 0 26             | 0 26       | 7                 | 0 26.  | 0   | 0 26.  | 0 26     | 0 26.3 | 9        | 0 26.3             | 0 26.3            | 30 26.3 1          |
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## ALIGNMENTS

```
APPLICANT: Dairiki, Kazuo
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Dise
FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT FILING DATE: 2997-00-10
PRIOR FILING DATE: 1997-00-10
PRIOR APPLICATION NUMBER: US 09/142,524
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PRIOR FILING DATE: 1998-09-09
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; COTHER INFORMATION: Cryjl peptide, Figure 1, Row 49
(S-10-354-240-63
Sequence 63, Application US/10354240; Publication No. US20030185847A1; GENERAL INFORMATION:
APPLICANT: Sone, Toshio; APPLICANT: Rume, Akinori
   TYPE: PRT
ORGANISM: Cryptomeria japonica
  NAME/KEY: MISC_FEATURE
   SEQ ID NO 63
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Gaps

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Sequence 65, Application US/10354240

Sequence 65, Application No. US200018847A1

GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Rume, Akinori
APPLICANT: Mama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: POST IN UNDER: US/10/354,240
CURRENT APPLICATION NUMBER: US/097/00740
PRIOR PLIING DATE: 1997-03-10
PRIOR PLIING DATE: 1997-03-10
PRIOR PLIING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
  APPLICANT: FOWLKES, Dana M.
APPLICANT: BARNETT, Thomas R.
APPLICANT: BARNETT, Thomas R.
APPLICANT: BARNETT, Thomas R.
APPLICANT: BURNETT, Thomas R.
APPLICANT: BURNETT, Thomas R.
APPLICANT: BURNETT, DANA BETHOD G. IDENTIFYING CONFORMATION-SENSITIVE BINDING PEPTIDES ANI TITLE OF INVENTION: THEREOF FILE REPERENCE: PAIGE-1H
CURRENT APPLICATION NUMBER: US 09/614,865
PRIOR APPLICATION NUMBER: US 09/614,865
PRIOR APPLICATION NUMBER: US 09/614,865
PRIOR APPLICATION NUMBER: US 09/614,865
NUMBER OF SEQ ID NOS: 268
NUMBER OF SEQ ID NOS: 268
SOFTWARE: PATENTIN VERSION 3.1
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; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 51
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Pred. No.
   Sequence 100, Application US/10346162
Publication No. US20030224390A1
GENERAL INFORMATION:
APPLICANT: KARO BIO USA, INC.
APPLICANT: FOWLKES, Dana M.
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100.0%; Pre
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   ò
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  APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Bairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Mama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103D1
CURRENT FILING DATE: 103-10
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
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LENGER PATENTIN VERSION 3.1
SEQ ID NO 64
  TITILE OF INVENTION NUMBER: US/10/354,240
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CURRENT PILLING DATE: 10303-01-29
PRIOR PILLING DATE: 1997-03-10
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PRIOR PILLING DATE: 1997-03-10
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PRIOR PILLING DATE: 1998-09-09
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Pred. No. 1.4e-05;
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   Sequence 62, Application US/10354240
Publication No. US20030185847A1
GENERAL INFORMATION:
APPLICANT: Some, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
  Sequence 64, Application US/10354240 Publication No. US20030185847A1 GENERAL INFORMATION:
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Sequence 142, Application US/10346162
| Publication No. US20030224390A1
| Publication No. US20030224390A1
| GENERAL INFORMATION:
| APPLICANT: KARO BIO USA, INC.
| APPLICANT: FOWLKES, Dana M.
| APPLICANT: BURNETT, Thomas R.
| APPLICANT: BURNETT, DATE: DATE: US/10/346,162
| TITLE OF INVENTION: THEREOF
| TITLE OF INVENTION: THEREOF
| CURRENT PELLING DATE: 2003-01-17
| PRIOR APPLICATION NUMBER: US 09/614,865
| PRIOR APPLICATION NUMBER: US 09/614,865
| PRIOR APPLICATION NUMBER: US 09/610,628
| PRIOR APPLICATION DATE: 2000-07-12
| PRIOR APPLICATION NUMBER: US 09/606,688
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   32.5%; Score 37; DB 14; Length 16; 55.6%; Pred. No. 1.2e+02; ive 2; Mismatches 2; Indels
  Ouery Match 30.7%; Score 35; DB 14; Length 13; Best Local Similarity 71.4%; Pred. No. 2e+02; Matches 5; Conservative 1; Mismatches 1; Indels
   ; OTHER INFORMATION: Polycarbonate-binding peptide 69 US-10-300-694A-69
  APPLICANT: Grinstaff, Mark W.
APPLICANT: Grinstaff, Mark W.
APPLICANT: Renam, Daniel J.
APPLICANT: Renam, Daniel J.
APPLICANT: Middleton, Crystan
TITLE OF INVENTION: INTERFACIAL BIOMATERIALS
FILE REFERENCE: 180/143/2
CURRENT APPLICATION NUMBER: US/10/300,694A
   PRIOR APPLICATION NUMBER: 2003-05-07
PRIOR APPLICATION NUMBER: US 60/331,843
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 117
SOFTWARE: Patentin version 3.2
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; PRIOR FILING DATE: 1998-06-22
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Publication No. US20030044851A1

GENERAL INFORMATION:

APPLICANT: Ruben et al.

ITILE OF INVENTION: 94 Human Secreted Proteins
FILE REPERENCE: PZ029P1

CURRENT APPLICATION NUMBER: US/10/012.542

PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/461,325

PRIOR PILING DATE: EARLIER FILING DATE: 1999-12-14

PRIOR PILING DATE: EARLIER FILING DATE: 1999-12-14

PRIOR PILING DATE: EARLIER FILING DATE: 1999-06-16

PRIOR PELICATION NUMBER: EARLIER APPLICATION NUMBER: 60/089,509

PRIOR PELING DATE: EARLIER PILING DATE: 1998-06-16

PRIOR PPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/089,509

PRIOR PPLING DATE: EARLIER FILING DATE: 1998-06-16

PRIOR PPLING DATE: EARLIER PILING DATE: 1998-06-22

PRIOR PLING DATE: EARLIER PILING DATE: 1998-06-22

PRIOR PROFESTION NUMBER: EARLIER PILING DATE: 1998-06-22
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Pred. No. 86;
5; Mismatches 5; Indels
   PUDLICATION NO. US20030065151A1

PUDLICATION NO. US20030065151A1

GENERAL INFORMATION al.

TITLE OF INVENTION at Human Secreted Proteins

FILE REPRENCE: PSO29330451D2

CURRENT APPLICATION NUMBER: US/10/115,123

CURRENT FILING DATE: 2002-04-04

FRIOR PILION DATE: 1999-06-15

FRIOR PILING DATE: 1998-06-16

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FRIOR PILING DATE: 1998-06-16

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Publication No. US20040235026A1
Sequence 1274, Application US/10813638
Publication No. US20040235026A1
GENERAL INFORMATION:
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin D.
TITLE OF INVENTION: USE THEREOF
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TITLE OF INVENTION: USE THEREOF
TITLE REPERENCE: 15966-599
CURRENT APPLICATION NUMBER: US/10/813,638
CURRENT APPLICATION NUMBER: US/10/813,783
PRIOR APPLICATION NUMBER: 00/163,783
PRIOR PRILICATION NUMBER: 00/163,783
PRIOR PRILICATION DATE: 1999-11-24
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US-10-184-9364-191
Sequence 191, Application US/10182936A
Sequence 191, Application No. US20040038860A1
GENERAL INFORMATION:
APPLICANT: Allen, Kristina M.
APPLICANT: Allen, Kristina M.
APPLICANT: Bhat, Bheem
APPLICANT: Bhat, Bheem
APPLICANT: Bragners, Veronique
APPLICANT: Robinson, John
APPLICANT: Waworsky, Paul
TITLE OF INVENTION: Reagents and Method for Modulating DKK-Mediated Interactions
TITLE OF INVENTION: Reagents us (10/182,936A)
CURRENT FILING DATE: 2002-08-02
PRIOR PRILORATION NUMBER: US 60/291,311
PRIOR PILING DATE: 2002-05-17
PRIOR PILING DATE: 2002-05-17
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR PILING DATE: 2002-02-01
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PRIOR PILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 216
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57.1%; Pred. No. 4.4e+02;
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US-10-182-936A-191
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Sequence 26, Application US/10468543
Sequence 26, Application US/20040091938A1
GENERAL INFORMATION:
GAPLICANT: Irimura, Tateuro
APPLICANT: Vim, Mijung
APPLICANT: Vim, Mijung
APPLICANT: Vim, Mijung
APPLICANT: Ono, Takashi
TITLE OF INVENTION: Lectins for Analyzing Sugar Chains and Method of Using the Same FILE REFERENCE: 03-786
CURRENT APPLICATION NUMBER: US/10/468,543
CURRENT PILING DATE: 2003-08-20
PRIOR APPLICATION NUMBER: JP 2001-044221
PRIOR PILING DATE: 2003-08-20
NUMBER OF SEQ ID NOS: 95
NUMBER OF SEQ ID NOS: 95
NUMBER OF SEQ ID NOS: 95
   APPLICANT: Timura, Tateuro
APPLICANT: Wijung
APPLICANT: Yim, Mijung
APPLICANT: Ono, Takashi
TITLE OR: INVENTION: Lectins for Analyzing Sugar Chains and Method of Using the Same
FILE REFERENCE: 03-786
CURRENT APPLICATION NUMBER: US/10/468,543
CURRENT FILING DATE: 2003-08-20
PRIOR APPLICATION NUMBER: UP 2001-044221
PRIOR FILING DATE: 2001-02-20
NUMBER OF SEQ ID NOS: 95
SOFTWARE: Patentin version 3.1
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Pred. No. 2.3e+02;
3; Mismatches 3.
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   Sequence 275, Application US/10280066
| Publication No. US2030180718A1
| GENERAL INFORMATION:
| APPLICANT: Pillutla, Renuka C. |
| APPLICANT: Pillutla, Renuka C. |
| APPLICANT: Brissette, Renee |
| APPLICANT: Brissette, Renee |
| APPLICANT: Brissette, Michael |
| APPLICANT: Blume, Arthur J. |
| APPLICANT: Blume, Arth
  APPLICANT: Pillutla, Renuka C.
APPLICANT: Pillutla, Renuka C.
APPLICANT: Pillutla, Renee
APPLICANT: Brissette, Renee
APPLICANT: Brissette, Michael
APPLICANT: Blume, Arthur J.
APPLICANT: Blume, Arthur J.
APPLICANT: Prendergast, John
APPLICANT: Prendergast, John
TITLE OF INVENTION: TARGET SPECIFIC SCREENING AND ITS USE FOR IDENTIFYING TARGET BIND
FILE REFERENCE: 2588-44001831
CURRENT FILING DATE: 2002-10-24
PRIOR FILING DATE: 2001-10-24
PRIOR FILING DATE: 2001-10-24
NUMBER OF SEQ ID NOS: 537
SOFTWARE: Patentin version 3.1
SQ ID NO 276
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TYPE: PRI
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  ; FEATURE:
; NAME/KEY: MISC FEATURE
; OTHER INFORMATION: 0700902-Hras-20M-PP-BC-B6
US-10-280-066-275
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US-10-182-936A-191
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US-10-280-066-276
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100, App 108, App 80, Appl 8, Appli 8, Appli 16, Appl

Sequence Sequence Sequence Sequence

Sequence

Sequence Sequence Sequence

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US-08-467-023-50

i Sequence 50, Application US/08467023

patent No. 6090386

general information:
    APPLICANT Griffeth, Irwin J.;
    APPLICANT Bond, Julian F.;
    APPLICANT Garman, Richard D;
    APPLICANT Ruo, Mei-Chang;
    APPLICANT Ruor, Siu-mei H.;
    APPLICANT Seuge, Siu-mei H.;
    APPLICANT Seuge, Siu-mei H.;
    APPLICANT Steven Andrew;
    APPLICANT STEVEN ANDRESS:
    APPLICANT STEVEN ANDRESS:
    CORRESPONDENCE ADDRESS:
    ADDRESSE: Immulogic Pharmaceutical Corporation, Inc.
    STREET: MA
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION MUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSTFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKT NUMBER: 38,872
REFERENCE/DOCKT NUMBER: 025.6 USD2 (IMI-028CPD2)
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  TELEFAX: (617) 227-5941
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APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disertile Reference: 870-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: D97-09-09
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
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   APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From
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   ImmuLogic Pharmaceutical Corporation, Inc.
   ; LOCATION: (1). (15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 48
US-09-142-24D-62
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'S Gouene 258, Application US/08467023
'Patent No. 6090386
   Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
  Yeung, Siu-mei H.,
Brauer, Andrew;
Exley, Mark A.;
  Garman, Richard D;
Kuo, Mei-Chang;
  TYPE: PRT
ORGANISM: Cryptomeria japonica
   71.9%;
   50.9%;
  6 DPWSIYAIGGSSNPT 20
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Best Local Similarity 93.3
Matches 14; Conservative
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  GENERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
   Query Match 50.5
Best Local Similarity 90.0
Matches 9; Conservative
   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
  1 ANNNYDPWSI 10
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   GENERAL INFORMATION:
APPLICANT: Griffe
  Waltham
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  ADDRESSEE:
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  8
  APPLICANT: Sone, Tochio
APPLICANT: Sone, Tochio
APPLICANT: Sone, Akinori
APPLICANT: Wame, Akinori
APPLICANT: Wame, Akinori
APPLICANT: Iwama, Akiko
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Disea
TITLE OF INVENTION: 1998-09-09
CURRENT APPLICATION NUMBER: US/09/142,524D
PRIOR APPLICATION NUMBER: POT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
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TYPE: PRT
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FULE APPLICATION WUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 174
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   1 ANNNYDPWSIYAIGG 15
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  APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Ivama, Akiko
APPLICANT: Kino, Kohsuke
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US-09-142-524D-65; Sequence 65, Application US/09142524D; Patent No. 6719976
   ; Sequence 51, Application US/08467023; Patent No. 6090386
  TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
   38,872
  TYPE: PRT
ORGANISM: Cryptomeria japonica
  Query Match 50.9%;
Best Local Similarity 90.0%;
Matches 9; Conservative
                                 NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,
REFERENCE/DOCKET NUMBER:
            ATTORNEY/AGENT INFORMATION:
  LENGTH: 20 amino acids
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TOPOLOGY: linear
  APPLICANT: Sone, Toshio
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   US-08-467-023-49
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APPLICANT:
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   Gaps
   APPLICANT: Griffeth, Irwin J.;
APPLICANT: Griffeth, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Yeung, Sil-mei H.;
APPLICANT: Yeung, Sil-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exteven F.;
APPLICANT: Brauer, Andrew;
APPLICANT: Waltham
COUNTRY: WAITHAM
ZIP: ANDRESS:
APPLICANT: WAITHAM
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  Score 58; DB 3; Length 19; Pred. No. 0.025;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INPORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/POCKET NUMBER: 38,872
REFERENCE/POCKET NUMBER: 38,872
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REFERENCE/POCKET NUMBER: 38,872
RELEPHONE: (617) 227-7400
TELLECOMMUNICATION INPORMATION:
TELLECOMMUNICATION INPORMATION:
TELLEPHONE: (617) 227-7400
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
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APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
  1; Mismatches
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NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,20
   REFERENCE/DOCKET NUMBER:
  TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
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  US-08-290-448A-55
  RESULT 10
US-08-290-448A-55
  TOPOLOGY:
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   APPLICANT: Rogers, Bruce
APPLICANT: Riapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Raffar, Thorunn
APPLICANT: Row, Met-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 66 State Street, suite 510
CITY: Boston
STATE: Massachusetts
                APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From
TITLE OF INVENTION: Appanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSE: ImmuLogic Pharmaceutical Corporation, Inc.
STREET: 610 Lincoln St
CITY: Waltham
  Score 53; DB 3; Length 20;
Pred. No. 0.15;
  0; Indels
   ZIP: 02154
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MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTANARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release:#1.0, Version #1.25
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  TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 51:
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LENGTH: 20 amino acids
   (617) 227-7400
Brauer, Andrew
  Query Match
Best Local Similarity 100.
Matches 10; Conservative
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: USA
  1 YAIGGSSNPT 10
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   USA
  US-08-290-448A-55
   STATE: MA
  COUNTRY:
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| Patent No. 5698204 |
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| GENERAL INPORMATION |
| APPLICANT: Refers, Bruce |
| APPLICANT: Rafner, Thorunn |
| APPLICANT: Rafner, Street, 93 |
| CORRESPONDENCE ADDRESS: ADDRESSE: LAHIVE & COCKFIELD |
| STREET: 60 State Street, suite 510 |
| CITY: Boston |
| CITY: Bo
   Score 50; DB 1; Length 17; Pred. No. 0.36;
   STATE: BOSCHUSELTS
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Parentln Recompatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentln Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/29,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: Max 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: Max 71, 1989
ATTOMENY/AGENT IMPORMATION:
APPLICATION NUMBER: US/08/290,448A
PILING DATE: August 15, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: MAY 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: MARCH 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. MANCHAGOULAS
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-018CN
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
```

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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
   43.9%;
   TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
  MOLECULE TYPE: peptide FRAGMENT TYPE: internal
  Conservative
   Massachusetts
   2 NNNYDPWSIYA 12
   7 NNNYDRWGTYA 17
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  Best Local Similarity
Matches 8; Conserv
 US-08-461-939B-55
  US-08-461-939B-55
  RESULT 13
US-08-464-000-55
   CITY: I
STATE:
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0
   18-08-175-069A-55
Sequence 55, Application US/08175069A
Sequence 55, Application US/08175069A
Patent No. 5776761
GENERAL INFORMATION:
APPLICANT: Rogers, Bruce
APPLICANT: Rafnar, Thorunn
APPLICANT: Ruo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEGUENCES: 93
CORRESPONDENCE ADDRESS:
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  Score 50; DB 1; Length 17; Pred. No. 0.36; 3; Indels 0; Mismatches 3; Indels
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  3; Indels
   3; Indels
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ADDIO TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/175,069A

FLING APPLICATION DATA:

APPLICATION NUMBER: US 07/529,951

FLING DATE: March 17, 1989

ATTORNEY/AGENT INFORMATION:

NAME: AMY E. Mandragouras

REGISTRATION NUMBER: 36,207

REFERENCE/DOCKET NUMBER: 36,207

REFERENCE/DOCKET NUMBER: 111-018DV

TELECOMMUNICATION NUMBER: 111-018DV
  0; Mismatches
  E: LAHIVE & COCKFIELD, LLP 60 State Street
   TELEFAX: (617)227-5941 INFORMATION FOR SEQ ID NO: 55:
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ilarity 72.7%;
Conservative (
   43.9%;
72.7%;
  (617) 227-7400
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LENGTH: 17 amino acids
17 amino acids
  8; Conservative
   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
  STREET: 60 State Stre
CITY: Boston
STATE: Massachusetts
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FRAGMENT TYPE: internal
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  2 NNNYDPWSIYA 12
   NNNYDRWGTYA 17
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   Query Match
Best Local Similarity
Matches 8; Conserv
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  Query Match
Best Local Similarity
Matches 8; Conserv
  TELEPHONE:
  ADDRESSEE:
  US-08-290-448A-55
   US-08-175-069A-55
                  TYPE: am
TOPOLOGY:
   TOPOLOGY:
LENGTH:
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   ò
   원
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RESULT 12

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Sequence 55, per late on US/084613958

CREMENAL INTOMACION:
APPLICANT: Rispec. David G.
APPLICANT: Rispec. Theory of G.
APPLICANT: Rispec. Theory of G.
APPLICANT: Rispec. Theory of G.
APPLICANT: Rispec. ADDRESS:
CONTRESS OF SEQUENCES:
CONTRESS OF SECUENCES:
CONTRESS OF SECUENCES.
CONTRESS OF SECUENCE
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   Sequence 10, Application US/09379665D
Patent No. 6509184
GENERAL INFORMATION
TITLE OF INVENTION: ALKALINE TOLERANT DEXTRANASE FROM STREPTOMYCES ANULATUS
FILLE REFERENCE: NREL 98-37
CURRENT APPLICATION NUMBER: US/09/379,665D
CURRENT FILING DATE: 1999-08-24
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APPLICATION NUMBER: US 07/529,951
FILING DATE: 29-MAY-1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: 17-MAR-1989
ATONNEY, AGENT INPORMATION:
NAME: Amy E. MANGEAGOUAE
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-018CN2
TELECHONE: (617)227-7400
   ZIP: 02109-1875
COMPUTER READABLE FORM:
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OPERATING SYSTEM: PC-DOS/MS-1
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SOFTWARE: PatentIn version 3.1
   TELEFAX: (617)227-5941
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60 State Street
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  Best Local Similarity 72.7
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RESULT 15

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Sequence 10. Application US/08861153A

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| Sequence 10. Application US/08861153A |
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| Patent No. 6723646 |
| Patent No. 6723604 |
| Patent No. 67236 |
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January 26, 2005, 16:03:29; Search time 55.4 Seconds (without alignments) 130.429 Million cell updates/sec
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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1 ILSEGNSFTAPNDSDKKEVT 20
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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|--------------|-----------------|------------------|------------------|------------------|------------------|-------------------|-------------------|------------------|-------------------|------------------|-------------------|-------------------|-------------------|-------------------|
|              |                 | 67, 1            | 68, 1            | 66, 1            | 69, 7            | 117,              | 60, 7             | 60, 7            | 31, 7             | 67, 1            | _                 | 145,              | 149,              | 153,              |
|              | Description     | Seguence 67,     | Seguence         | Sequence         | Sequence         | Sequence          | Sequence          | Sequence         | Sequence          | Seguence         | Sequence          | Sequence          | Sequence          | Sequence          |
| SUMMARIES    | 1D              | US-10-354-240-67 | US-10-354-240-68 | US-10-354-240-66 | US-10-354-240-69 | US-10-354-240-117 | US-09-966-459A-60 | US-10-323-412-60 | US-10-172-425B-31 | US-10-432-422-67 | US-10-632-706-141 | US-10-632-706-145 | US-10-632-706-149 | US-10-632-706-153 |
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| ,1<br>1<br>1 | watch bength UB | 15               | 15               | 15               | 15               | 15                | 15                | 15               | 14                | 15               | 17                | 17                | 17                | 17                |
| %<br>Query   | marcn           | 66.3             | 61.4             | 47.5             | 36.6             | 32.7              | 31.7              | 31.7             | 30.7              | 30.7             | 30.7              | 30.7              | 30.7              | 30.7              |
|              | score           | 67               | 62               | 48               | 37               | 33                | 32                | 32               | 31                | 31               | 31                | 31                | 31                | 31                |
| Result       |                 | П                | 7                | m                | 4                | S                 | 9                 | . 7              | 80                | 6                | 10                | 11                | 12                | 13                |

Gaps ö

Query Match 66.3%; Score 67; DB 14; Length 15; Best Local Similarity 92.9%; Pred. No. 0.00059; Matches 13; Conservative 1; Mismatches 0; Indels

1 ILSEGNSFTAPNDS 14

COCATION: (1).7(15)
COTHER INFORMATION: Cryjl peptide, Figure 1, Row 53
US-10-354-240-67

TYPE: PRT ORGANISM: Cryptomeria japonica

NAME/KEY: MISC\_FEATURE

FEATURE:

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|-----|----------------------------------------|--------|-----------------------------------|------------------|----------------|-------------------------------------------|---------------------------------------------------------------|
|     | 14                                     | 31     | m                                 | 17               | 16             | н,                                        | equence 157,                                                  |
|     | 15                                     | E 6    | ግ (                               | 200              | վ ,            | US-10-057-789-298                         | equence 298,                                                  |
|     | 9 F                                    | 7 0    | <b>n</b> (                        | 2 5              | <del>7</del> , | V                                         | , a                                                           |
|     | \ C                                    | 9 6    | 40                                | 1 1              | 1 -            | US-10-1/2-4235-44<br>HS-10-777-893-65     | Sequence 44, Appl                                             |
|     | 19                                     | 8 8    | 10                                | 15               | , 0            | US-09-739-852-8                           | Sequence 8. Appli                                             |
|     | 20                                     | 29     |                                   | 15               | 14             | US-10-354-240-116                         | Sequence                                                      |
|     | 21                                     | 59     | ~                                 | 15               | w              | US-10-203-915A-126                        | Sequence 126, App                                             |
|     | 22                                     | 53     | 7                                 | 16               | δ.             | US-09-889-468-44                          | Sequence 44, Appl                                             |
|     | 23                                     | 50     | 7                                 | 17               | 77             | US-10-160-232-16                          | ednen                                                         |
|     | 27 c                                   | 0 0    | C) (                              | 17               | 4.             | US-10-281-479A-26                         | ednen                                                         |
|     | 2 K                                    | N (    | 7                                 | 7:               | 4.             | US-10-2/5-18UA-26                         | equen                                                         |
|     | 9 10                                   | 2 0    | 710                               | 1.               | 4 1            | US-IU-286-132A-26                         | Sequence 26, Appl                                             |
|     | 7 00                                   | N C    | <b>7</b> C                        | - a              | 1              | 11C-10-4//-3/19                           | manha                                                         |
|     | 0 0                                    | A C    | 4 C                               | 9 6              | # <u>~</u>     | 175-10-004-013-202                        | manha                                                         |
|     | V C                                    | 9 0    | 4 (                               | 9 6              | 7 7            | US-10-084-813-428                         | משושט                                                         |
|     | , r                                    | 9 0    | 10                                | 9 6              | 7 7            | TIS-10-084-813-420                        | Semience 430. App                                             |
|     | 100                                    | 9 0    | 10                                | 9 6              | 7              | TIS-10-084-813-431                        |                                                               |
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|     | 7 6                                    | 0 0    | 4 C                               | 9 6              | * 4            | 115-10-034-40/A-13                        | Sequence 19, App.                                             |
|     | # U                                    | 200    | 4 C                               | 7 7              | ט ט            | 113-10-303-330-17                         | Sequence 17, Appr                                             |
|     | 7 7                                    |        | 4 C                               | † C              | 10             | 116-10-020-116-10-01                      | Company 22 April                                              |
|     | 0 0                                    | 0 0    | 4 C                               | 0 0              | ·) u           | US-10-000-100-23                          |                                                               |
|     | 2 0                                    | 0 0    | 4 (                               | ים<br>ד          | 2              | 7 .                                       | מהליה זה שלהיה להיה אלה היה שלהיה שלה להיה להיה להיה להיה להי |
|     | 200                                    | 0 0    | 71 (                              | ς ;              | , ,            | US-US-966-900A-32                         | Sequence 32, Appr                                             |
|     | 5.5                                    | 80 0   | ~ (                               | Ţ;               | 2 :            | ຸດ                                        | Sequence 52, Appl                                             |
|     | 0 .                                    | 8 6    | 71 (                              | 1.               | 9 ,            | 5-09-880-748-2                            |                                                               |
|     | 14.                                    | 80 6   | ~ (                               | Υ ·              | -1 +<br>4 t    | US-10-293-418-2991                        | o ·                                                           |
|     | 42                                     | 128    |                                   | 15               | 5              | S-10-442-909-4                            | Sequence                                                      |
|     | 43                                     | 28     | (7)                               | 16               | σ,             | -09-757-774                               | equence                                                       |
|     | 44                                     | 9 0    | 27                                | 9 4              | ט י            | US-09-757-77                              | quence                                                        |
|     | <b>.</b>                               | 8      | 7                                 | D<br>T           | n              | US-10-449-829A                            | riddw 'c acinnbac                                             |
|     |                                        |        |                                   |                  |                | ALIGNMENTS                                |                                                               |
|     |                                        |        |                                   |                  |                |                                           |                                                               |
| -   | RESULT 1                               |        |                                   |                  |                |                                           |                                                               |
| _   | US-10-354-240-67                       | -240-6 |                                   |                  |                |                                           |                                                               |
|     | ; Sequence 67, Application US/10354240 | 9 67,  | Applica                           | tion US          | 7105           | 354240                                    |                                                               |
|     | CENEDAL INFORMATION                    | TNEOD  | WATTON.                           | 007070           | ć              | -1                                        |                                                               |
|     | GENERAL<br>TODAL TO                    | MT.    | ADDITONT CODE TORY                | o t do           |                |                                           |                                                               |
|     | . ADDLICANT.                           | . EN   | Kume akinori                      | kinori           |                |                                           |                                                               |
|     | APPLICANT                              |        | Dairiki                           | . Kazuo          | _              |                                           |                                                               |
|     | APPLICANT                              |        | Iwama.                            | Akiko            |                |                                           |                                                               |
|     | APPLICANT                              | INT:   | Kino, Kohsuke                     | ohsuke           |                | ,                                         |                                                               |
|     | ; TITLE                                | OF INV | TITLE OF INVENTION: Peptide-Based | Peptid           | le-Ba          | ased Immunotherapeutic Agent              | nt for Treating Allergic Dise                                 |
| Ī   | , FILE RI                              | SPEREN | ICE: SPO                          | -103D1           |                |                                           | •                                                             |
|     | ; CURRENT                              | r APPL | ICATION                           | NUMBER           | : UE           | CURRENT APPLICATION NUMBER: US/10/354,240 |                                                               |
|     | ; CURRENT                              | r FILI | CURRENT FILING DATE: 2003-01-2    | : 2003           | -01-           | -29                                       |                                                               |
| -   | , PRIOR !                              | APPLIC | ATION N                           | UMBER:           | PCT)           | PRIOR APPLICATION NUMBER: PCT/JP97/00740  |                                                               |
|     | PRIOR F                                | TILING | FILING DATE: 1997-03-10           | 1997-03          | 9              |                                           |                                                               |
| -   | PRIOR                                  | APPLIC | PRIOR APPLICATION NUMBER: US      | UMBER:           | 3              | 09/142,524                                |                                                               |
| - ' | NIMBER                                 | TO TO  | ON CIT OF                         | 1330-03<br>S 174 | ,<br>,         |                                           |                                                               |
|     | SOFTWAR                                | E: Pa  | SOFTWARE: PatentIn version        | version          | 3.1            |                                           |                                                               |
|     | SEQ ID NO                              | 70 67  |                                   | 1                |                |                                           |                                                               |
|     | ; LENGTH:                              |        |                                   |                  |                |                                           |                                                               |
|     | TVDE . DDT                             | - 00   |                                   |                  |                |                                           |                                                               |

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TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Dise-
FILE REFERENCE: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT FILING DATE: 2003-01-29
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1997-03-10
PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 69
   TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disercine Reference: SPO-103D1
CURRENT APPLICATION NUMBER: US/10/354,240
CURRENT FILING DATE: 2003-01-29
PRIOR APPLICATION NUMBER: PCT/JP97/00740
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PRIOR APPLICATION DATE: 1999-09-09
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   Score 37; DB 14; Length 15;
Pred. No. 46;
  Best Local Similarity 100.0%; Pred. No. 0.74;
Matches 10; Conservative 0; Mismatches 0; Indels
   ; LOCATION: (1). (15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 55
US-10-354-240-69
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  ; Sequence 69, Application US/10354240; Publication No. US20030185847A1; GENERAL INFORMATION:
; APPLICANT: Sone, Toshio; APPLICANT: Rume, Akinori; APPLICANT: Dairiki, Kazuo; APPLICANT: Iwama, Akiko; APPLICANT: Iwama, Akiko; APPLICANT: Kino, Kohsuke
  Sequence 117, Application US/10354240
Publication No. US20030185847A1
GENERAL INFORMATION:
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70.0%;
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APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
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US-10-354-240-117
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CURRENT FILING DATE: 2003-01-29
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PRIOR FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 174
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    MS-10-354-240-68

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OTHER INFORMATION: Cryjl peptide, Figure 1, Row
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   Sequence 66, Application US/10354240
Publication No. US20030185847A1
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Publication No. US20030185847A1
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  ORGANISM: Cryptomeria japonica
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   NSFTAPNESYKKOVT 15
1 ILSEGNSFTAPNES 14
   Kume, Akinori
Dairiki, Kazuo
Iwama, Akiko
Kino, Kohsuke
   Query Match
Best Local Similarity 80.0
Matches 12; Conservative
   APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
  APPLICANT: Sone, Toshio
   NAME/KEY: MISC FEATURE
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APPLICANT:
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Gaps
   Gaps
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  JAPELICANT: Kaempfer, Raymond

APPLICANT: Arad, Gila

APPLICANT: Arad, Gila

TITLE OF INVENTION: BROAD SPECTRUM ANTAGONISTS AND VACCINES

TITLE OF INVENTION: DIRECTED AGAINST PYROGENIC EXOTOXINS

FILE REFERENCE: A31967-PCT-USA-A-A 066031.0164

CURRENT APPLICATION NUMBER: US/10/172,425B

PRIOR FILING DATE: 1998-09-10

PRIOR APPLICATION NUMBER: 09/150,947

PRIOR PILING DATE: 1998-09-10

PRIOR PILING DATE: 1997-12-30

PRIOR FILING DATE: 1997-12-30

NUMBER OF SEQ ID NOS: 57

SOFTWARE: FastsEQ for Windows Version 4.0

LENGTH: 14

TYPE: PRI
  Score 32; DB 16; Length 15;
Pred. No. 3e+02;
1; Mismatches 4; Indels
   30.7%; Score 31; DB 14; Length 14; larity 85.7%; Pred. No. 4e+02; Conservative 1; Mismatches 0; Indels
   APPLICANT: Syngenta Participations AG
APPLICANT: Cornell Research Foundation, Inc.
APPLICANT: Yoder, Olen
APPLICANT: Turgeon, Barbara G.
APPLICANT: Lu, Shen-wen
TITLE OF INVENTION: Fungal Iron Reductase Gene
FILE REFERENCE: 1360.017W01
CURRENT APPLICATION NUMBER: US/10/432,422
  FEATURE:
; OTHER INFORMATION: Synthesized Polypeptide.
US-10-323-412-60
              APPLICATION NUMBER: U.S. 60/261,776
   US-10-172-425B-31
, Sequence 31, Application US/10172425B
; Publication No. US20030147908A1
; GENERAL INFORMATION:
   ; Sequence 67, Application US/10432422; Publication No. US20040076981A1; GENERAL INFORMATION:
PRIOR APPLICATION NUMBER: U.S. V., PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: U.S. 60, PRIOR FILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-08-17
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin version 3.1
SEQ ID NO 60
  ORGANISM: Streptococcus pyogenes
   TYPE: PRT
ORGANISM: Artificial Sequence
  h 31.7%;
Similarity 54.5%;
6; Conservative
   3 SDGESFFEPGD 13
  3 SEGNSFTAPND 13
  Query Match
Best Local Similarity
Matches 6; Conserval
   14 SDKKEVT 20
  Best Local Similarity
Matches 6; Conserv
  3 TDKKEVT 9
   US-10-172-425B-31
  US-10-432-422-67
   g
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  셤
   APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY4, AND METHODS OF
TITLE OF INVENTION: THEREOF
TITLE OF INVENTION: THEREOF
FILE REFERENCE: D0039A LTP
CURRENT APPLICATION NUMBER: US,10/323,412
CURRENT APPLICATION NUMBER: U.S. 09/964,459
PRIOR FILING DATE: 2001-2.6
PRIOR FILING DATE: 2001-2.6
PRIOR FILING DATE: 2000-0-2.6
PRIOR FILING DATE: 2000-09-27
   ó
   APPLICANT: BARBER, L.

APPLICANT: BARBER, L.

APPLICANT: RORNACKER, M.G.

TITLE OF INVENTION: A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY4,

TITLE OF INVENTION: AND LUNG

FILE REPERENCE: D0033NP

CURRENT APPLICATION NUMBER: US/09/966,459A

CURRENT APPLICATION NUMBER: 60/235,833

PRIOR PRICH DATE: 2000-09-26

PRIOR FILING DATE: 2000-09-27

PRIOR FILING DATE: 2001-01-16

PRIOR FILING DATE: 2001-01-16

PRIOR FILING DATE: 2001-07-13

PRIOR FILING DATE: 2001-07-13

PRIOR FILING DATE: 2001-07-13

PRIOR FILING DATE: 2001-07-13

PRIOR FILING DATE: 2001-08-17

NUMBER OF SEQ ID NOS: 60

SOFTWARE: Patentin Ver. 2.1
   Gaps
   Gaps
   OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: polypeptide
   ö
   .;
0
  32.7%; Score 33; DB 14; Length 15; 70.0%; Pred. No. 2.1e+02; Live 0; Mismatches 3; Indels
  31.7%; Score 32; DB 10; Length 15; 54.5%; Pred. No. 3e+02; tive 1; Mismatches 4; Indels
   ; LOCATION: (1)..(15)
; OTHER INFORMATION: Cryj2 peptide, Figure 2, Row 34
US-10-354-240-117
  Sequence 60, Application US/09966459A Publication No. US20030022237Al GENERAL INFORMATION:
  Sequence 60, Application US/10323412
Publication No. US20040121330A1
GENERAL INFORMATION:
   Jequence
Publication NC.
GENERAL INFORMATION:
APPLICANT: FEDER, J.N.
APPLICANT: RAMANATHAN, C.S.
APPLICANT: HAWKEN, D.R.
APPLICANT: HAWKEN, D.R.
APPLICANT: APPLICANT
  TYPE: PRT
ORGANISM: Artificial Sequence
  Query Match
Best Local Similarity 70.0 Matches 7; Conservative
  Query Match 31.7
Best Local Similarity 54.5
Matches 6; Conservative
  3 SEGNSFTAPND 13
  3 SDGESFFEPGD 13
   5 GNSFTAPNDS 14
  2 GISITAPRDS 11
  US-09-966-459A-60
   US-09-966-459A-60
  -10-323-412-60
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Sequence 153, Application US/10632706

Sequence 153, Application US/10632706

Publication No. US20040175385A1

GENERAL INFORMATION:

APPLICANT: MARKES, DAMES D.

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

TITLE OF INVENTION: BUBMOTOXINS

TITLE OF INVENTION: 40951200S

CURRENT APPLICATION NUMBER: US 60/400, 721

PRIOR APPLICATION NUMBER: US 60/400, 721

PRIOR FILING DATE: 1998-08-31

NUMBER OF SEQ ID NOS: 278

SOFTWARE: PATENT PA
  Sequence 149, Application US/10632706

Publication No. US20040175385A1

GENERAL INPORMATION:

APPLICANT: MARKS, JAMES D.

APPLICANT: AMERSDORFER, PETER

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

TITLE OF INVENTION: NEUROTOXINS

TITLE OF INVENTION: NEUROTOXINS

FILE REFERENCE: 4077-895120US

CURRENT APPLICATION NUMBER: US/10/632,706

CURRENT FILING DATE: 2002-08-01

PRIOR APPLICATION NUMBER: US 09/144,806

SOFTWARE: PALENTIN POST: 278
  Gaps
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  ;
0
  Score 31; DB 16; Length 17;
Pred. No. 5e+02;
   Score 31; DB 16; Length 17;
Pred. No. 5e+02;
  5; Indels
  ; FEATURE:
; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-145
   ; OTHER INFORMATION: single chain antibody fragment US-10-632-706-149
  Mismatches
  3; Mismatches
                                   SOFTWARE: PatentIn version 3.2
SEQ ID NO 145
LENGTH: 17
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  Query Match 30.7%;
Best Local Similarity 46.7%;
Matches 7; Conservative
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   :|:| |:| | | | | | | 2 ISDGGSYTYYPDSVK 16
  2 LSEGNSFTAPNDSDK 16
   2 LSEGNSFTAPNDSDK 16
   Query Match
Best Local Similarity 46.7
Matches 7; Conservative
       NUMBER OF SEQ ID NOS: 278
  TYPE: PRT
ORGANISM: Artificial
   TYPE: PRT
ORGANISM: Artificial
  RESULT 12
US-10-632-706-149
  RESULT 13
US-10-632-706-153
   FEATURE:
   셤
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  ð
   APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THEARDEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROPOXINS
FILE REFERENCE: 4077-895120US
FILE REFERENCE: 4077-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE PATENTING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
   Sequence 145. Application US/10632706
BUDICALION NO. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSONERS, PETER
APPLICANTI AMERSONERS, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REPERBNCE: 407T-895120US
CURRENT PAPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR FILING DATE: 2003-08-01
PRIOR FILING DATE: 3002-08-01
PRIOR FILING DATE: 3002-08-01
PRIOR FILING DATE: 3003-08-01
PRIOR FILING DATE: 3003-08-01
   ö
  ö
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   Query Match
30.7%; Score 31; DB 16; Length 17;
Best Local Similarity 46.7%; Pred. No. 5e+02;
Matches 7; Conservative 3; Mismatches 5; Indels
   Length 15;
  Indels
  , OTHER INFORMATION: single chain antibody fragment
US-10-632-706-141
   30.7%; Score 31; DB 15; 40.0%; Pred. No. 4.4e+02;
  4; Mismatches
CURRENT FILING DATE: 2003-05-21
PRIOR APPLICATION NUMBER: US 60/252,732
PRIOR FILING DATE: 2000-11-22
PRIOR APPLICATION NUMBER: US 60/252,649
PRIOR FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 210
SOFTWARE: PREESEQ for Windows Version 4.0
  Sequence 141, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
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2 ISDGGSYTYYPDSVK 16
   1 ILSEGNSFTAPNDSD 15
   : | |::|| : ||
1 VLKAGHAFTLIDPSD 15
   2 LSEGNSFTAPNDSDK 16
  TYPE: PRT
ORGANISM: Fusarium scirpi
   ORGANISM: Artificial
  US-10-632-706-145
  US-10-632-706-141
  US-10-432-422-67
   SEQ ID NO 141
LENGTH: 17
  SEQ ID NO 67
LENGTH: 15
  FEATURE:
   ò
   ò
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  엄
   APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
APPLICANT: MARKSDORFER, PETER
TITLE OF INVENTION: THEAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 4077-8951200S
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT PILING DATE: 2003-08-01
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOSTWARE PATENTING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
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0
   Sequence 298, Application US/10057789

Sequence 298, Application US/10057789

Publication No. US20030082522A1

GENERAL INFORMATION:

APPLICANT: Bull Haynes

APPLICANT: Join Wais

APPLICANT: John Yates

APPLICANT: John Yates

TITLE OF INVENTION: DIFFERENTIAL LABELING FOR QUANTITATIVE

TITLE OF INVENTION: DIFFERENTIAL LABELING FOR QUANTITATIVE

TITLE OF INVENTION: DAMALYSIS OF COMPLEX PROTEIN MIXTURES

FILE REFERENCE: NADI: 0.22A

CURRENT FILING DATE: 2002-06-28

PRIOR APPLICATION NUMBER: US 60/264,576

PRIOR APPLICATION NUMBER: US 60/264,576

PRIOR APPLICATION NUMBER: US 60/305,232

PRIOR FILING DATE: 2001-07-13

NUMBER OF SEQ ID NOS: 311

SOFTWARE: FBSESQ for Windows Version 4.0

SEQ ID NO 298

LENGTH: 20
   30.7%; Score 31; DB 16; Length 17; 46.7%; Pred. No. 5e+02;
  30.7%; Score 31; DB 16; Length 17; 46.7%; Pred. No. 5e+02; tive 3; Mismatches 5; Indels
  5; Indels
   ; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-157
  ; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-153
  3; Mismatches
  Sequence 157, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
  2 LSEGNSFTAPNDSDK 16
   2 LSEGNSFTAPNDSDK 16
  Query Match
Best Local Similarity 46.7<sup>3</sup>
Matches 7; Conservative
  Query Match
Best Local Similarity 46.7
Matches 7; Conservative
TYPE: PRT
ORGANISM: Artificial
  ORGANISM: Artificial
   RESULT 14
US-10-632-706-157
  SEQ ID NO 157
  FEATURE:
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   셤
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PERTURE:

| PERTURE:
| FEATURE:
| FEATURE:
| NAME/KEY: VARIANT |
| LOCATION: 4 |
| OTHER INFORMATION: Xaa = Modified Cysteine
US-10-057-789-298 | 30.7%; Score 31; DB 14; Length 20;
Best Local Similarity 50.0%; Pred. No. 6.18+02;
Matches 7; Conservative 0; Mismatches 7; Indels 0; Gaps

Qy 2 LSEGNSFTARNDSD 15
| | | | | | | |
| Db 1 LSYXGGLPAPEDSD 14
| Search completed: January 26, 2005, 16:55:06
Job time: 56.5 secs
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LENGTH: 20 amino acids TYPE: amino acid
   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
   TYPE: amino acid
TOPOLOGY: linear
  STREET: 610 --
USA
   ZIP: 02154
   ADDRESSEE:
  US-08-467-023-52
  STATE: M. COUNTRY:
Appli
Appli
   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
   January 26, 2005, 15:54:02 ; Search time 14.6 Seconds (without alignments) 90.847 Million cell updates/sec
   Sequence 113, Sequence 39, A Sequence 7, App Sequence 7, App Sequence 7, App Sequence 8, App
   Sequence 170,
Sequence 170,
Sequence 170,
Sequence 170,
Sequence 170,
Sequence 170,
  Sequence 52,
  Description
  Sequence 5
Sequence 6
Sequence 5
Sequence 3
   Sequence 1
Sequence 2
Sequence 3
Sequence 3
Sequence 1
Sequence 1
  Sequence (
   Sequence
  Issued Patents AA:*

('GgnZ_6')ptodata'/,iaa/5A_COMB.pep:*

('GgnZ_6')ptodata'/,iaa/5B_COMB.pep:*

('GgnZ_6')ptodata'/,iaa/6A_COMB.pep:*

('GgnZ_6')ptodata'/,iaa/6B_COMB.pep:*

('GgnZ_6')ptodata'/,iaa/PCTUS_COMB.pep:*

('GgnZ_6')ptodata'/,iaa/PCTUS_COMB.pep:*
        GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
  US-09-142-524D-68
US-09-142-524D-68
US-09-142-524D-68
US-08-467-023-51
US-08-467-023-53
US-09-014-416-33
US-09-014-416-35
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US-09-014-416-36
US-08-473-170
US-08-473-170
US-08-475-263-170
US-08-477-134-170
US-08-485-695-170
   US-08-295-643-15
US-08-804-439A-113
US-09-014-416-39
US-08-475-844-7
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  US-09-133-341-8
   478139 segs, 66318000 residues
  SUMMARIES
   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
  - protein search, using sw model
  US-09-202-464-29
101
1 ILSEGNSFTAPNDSDKKEVT 20
  BLOSUM62
Gapop 10.0 , Gapext 0.5
  Length
  Minimum DB seq length: 0 Maximum DB seq length: 20
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Perfect score:
   Scoring table:
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   OM protein
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   Searched:
   Database
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No.
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Sequence 29, Appl
Sequence 15, Appl
Sequence 50, Appl
Sequence 51, Appl
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Appli
  Sequence
Sequence
Sequence
Sequence
Sequence
Sequence
Sequence
Sequence
         Sequence 1
Sequence 4
Patent No.
  APPLICANT: Pollock, Joanne;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
APPLICANT: Garman, Richard D;
APPLICANT: Kuo, Mel-Chang;
APPLICANT: Kuo, Mel-Chang;
APPLICANT: Young, Siu-mei H.;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM:
COMPUTER: IBM PC compatible
OPERATING SYSTEM:
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PRICATION NUMBER: US/08/467,023
FLIGHTON NUMBER: US/08/467,023
FLIGHTON NUMBER: US/08/467,023
FLIGHTON NUMBER: US/25
FLIGHTON NUMBER: 08/350,225
FLIGHTON NUMBER: 08/350,225
FLIGHTON NUMBER: 08/350,225
FLIGHTON NUMBER: 38,872
REFERRENCE/DOCKET NUMBER: 38,872
REFERRENCE/DOCKET NUMBER: 38,872
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHOMMUNICATION INFORMATION:
TELECHONE: (617) 227-7400
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTER/STICS:
TENACH. 20 amino acids
  3: ImmuLogic Pharmaceutical Corporation, Inc 610 Lincoln St
US-09-142-5245-116
US-09-014-416-42
US-08-18-252-29
US-08-218-025A-15
PCT-US93-11703-51
US-08-440-322-3
US-08-440-331-4
US-08-440-331-4
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US-08-448-252-26
US-08-448-252-26
US-08-448-252-26
US-08-448-253-26
US-08-448-253-26
US-08-448-85-86-190-23
US-08-448-85-86-157
US-08-445-86-157
   ALIGNMENTS
   Sequence 52, Application US/08467023
Patent No. 6090386
GENERAL INFORMATION:
```

```
APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Iwana, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR FILING DATE: 1997-03-10
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
SEQ ID NO 66
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  Gaps
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0
  TITLE OF INVENTION: Allergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
  Length 15;
   Length 15;
  0; Indels
  ADDRESSEE: ImmuLogic Pharmaceutical Corporation, Inc. STREET: 610 Lincoln St
   Score 62; DB 4; Ler
Pred. No. 0.0011;
    OTHER INFORMATION: Cryjl peptide, Figure 1, Row 54
  ; LOCATION: (1)...(15)
; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 52
VGS-09-142-524D-66
   Score 48; DB 4;
Pred. No. 0.21;
   61.4%; Scor.
80.0%; Pred. No. u.v.
2; Mismatches
   47.5%; Scor.
100.0%; Pred. No. ...
   Sequence 66, Application US/09142524D Patent No. 6719976
GENERAL INFORMATION:
APPLICANT: Sone, Toshio APPLICANT: Kume, Akinori APPLICANT: Dairiki, Kazuo
   Sequence 51, Application US/08467023 Patent No. 6090386
  APPLICANT: Garman, Richard D.
APPLICANT: Kuo, Mei-Chang;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Exley, Mark A.;
APPLICANT: Exley, Mark A.;
APPLICANT: Powers, Steven P.
   Griffeth, Irwin J.;
Pollock, Joanne;
Bond, Julian F.;
   TYPE: PRT
ORGANISM: Cryptomeria japonica
  1 NSFTAPNESYKKQVT 15
   6 NSFTAPNDSDKKEVT 20
   Query Match
Best Local Similarity 80.0
Matches 12; Conservative
   Query Match
Best Local Similarity 100.
Matches 10; Conservative
  CORRESPONDENCE ADDRESS:
ADDRESSEE: ImmuLogic
   1 ILSEGNSFTA 10
   6 ILSEGNSFTA 15
   NAME/KEY: MISC_FEATURE
  NUMBER OF SEQUENCES:
  GENERAL INFORMATION:
APPLICANT: Griffer
APPLICANT: Pollocl
  Waltham
   RESULT 4
US-09-142-524D-66
         ; US-09-142-524D-68
  LOCATION:
   g
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  APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Dairiki, Kazuo
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohauke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REPERBUES: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
   GENERAL INFORMATION:

APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Twama, Akinori
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Iwama, Akiko
APPLICANT: Kino, Kohsuke
TILE REPERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT APPLICATION NUMBER: DS/09/192
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR PILING DATE: 1997-03-10
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  Gaps
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0
  84.2%; Score 85; DB 3; Length 20; 85.0%; Pred. No. 2.9e-07;
  Query Match

66.3%; Score 67; DB 4; Length 15;

Best Local Similarity 92.9%; Pred. No. 0.00017;

Matches 13; Conservative 1; Mismatches 0; Indels
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; OTHER INFORMATION: Cryjl peptide, Figure 1, Row 53
US-09-142-524D-67
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  Sequence 67, Application US/09142524D Patent No. 6719976
  Sequence 68, Application US/09142524D Patent No. 6719976
  1 ILSEGNSFTAPNDSDKKEVT 20
   1 ILSEGNSFTAPNESYKKQVT 20
   ORGANISM: Cryptomeria japonica
   ORGANISM: Cryptomeria japonica
  NUMBER OF SEQ ID NOS: 174
SOFWRARE: Patentin version 3.1
SEQ ID NO 8
ENGTH: 15
  1 ILSEGNSFTAPNES 14
  1 ILSEGNSFTAPNDS 14
  Best Local Similarity 85.0 Matches 17; Conservative
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   FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(15)
  US-09-142-524D-67
US-08-467-023-52
  SEQ ID NO 67
LENGTH: 15
  Query Match
   TYPE: PRT
  ò
```

```
APPLICANT: Yanagi, Masayuki
APPLICANT: Bukh, Jens
APPLICANT: Bukh, Jens
APPLICANT: Emerson, Susanne U.
APPLICANT: Purcell, Robert H.
TITLE OF INVENTION: CLONED GENOMES OF INFECTIOUS HEPATITIS C VIRUSES AND
TITLE OF INVENTION: USES THEREOF
  Gaps
  .
0
  APPLICANT: Garman, Richard D,
APPLICANT: Kuo, Mei-Chang,
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Yeung, Siu-mei H.;
APPLICANT: Brauer, Andrew;
APPLICANT: Brauer, Andrew;
APPLICANT: Powers, Steven P.
TITLE OF INVENTION: Albergenic Proteins And Peptides From TITLE OF INVENTION: Japanese Cedar Pollen
NUMBER OF SEQUENCES: 261
  DB 3; Length 20;
  1; Indels
  CORRESPONDENCE ADDRESS:
ADDRESSE: ImmuLogic Pharmaceutical Corporation, Inc. STREET: 61 Lincoln St CITY: Waltham
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: June 6, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,892
REFERENCE/DOCKET NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (617) 227-7400
TELEFRAX: (617) 227-7400
TELEFRAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
   025.6 USD2 (IMI-028CPD2)
  36.6%; Score 37; DB 70.0%; Pred. No. 18; tive 2; Mismatches
  US-09-014-416-33
; Sequence 33, Application US/09014416
; Patent No. 6153421
; GENERAL INFORMATION:
   Application US/08467023
  GENERAL INFORMATION:
APPLICANT: Griffeth, Irwin J.;
APPLICANT: Pollock, Joanne;
APPLICANT: Bond, Julian F.;
   Garman, Richard D;
Kuo, Mei-Chang;
  Ouery Match
Best Local Similarity 70.0.
  MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-467-023-53
  ZIP: 02154
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  11 PNDSDKKEVT 20
  1 PNESYKKOVT 10
   linear
   USA
  COUNTRY: UR
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   PREERAL INFORMATION:
APPLICANT: Sone, Toshio
APPLICANT: Kume, Akinori
APPLICANT: Inama, Akinori
APPLICANT: Inama, Akino
APPLICANT: Inama, Akino
APPLICANT: Kino, Kohsuke
APPLICANT: Kino, Kohsuke
APPLICANT: Kino, Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
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0
  47.5%; Score 48; DB 3; Length 20; 100.0%; Pred. No. 0.3; tive 0; Mismatches 0; Indels
  36.6%; Score 37; DB 4; Length 15; 70.0%; Pred. No. 13;
   1; Indels
                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,023
FILING DATE: Unne 6, 1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/350,225
FILING DATE: December 6, 1994
ATTORNEY/ARTY INFORMATION:
NAME: Jane E. Remillard
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 38,872
REGISTRATION NUMBER: 025.6 USD2 (IMI-028CPD2)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
  LOCATION: (1). (15)
OTHER INFORMATION: Cryjl peptide, Figure 1, Row 55
  2; Mismatches
   FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
CURRENT FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: PCT/JD97/00740
PRIOR FILING DATE: 1997-03-10
SUMBER OF SEC ID NOS: 174
SOFTWARE: Patentin version 3.1
   Sequence 69, Application US/09142524D Patent No. 6719976
   ORGANISM: Cryptomeria japonica
   TELEPHONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 51
SEQUENCE CHARACTERISTICS:
1.ENGTH: 20 amino acids
  Query Match
Best Local Similarity 70.0
Matches 7; Conservative
  Query Match
Best Local Similarity 100.
Matches 10; Conservative
  MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
COMPUTER READABLE FORM:
  11 PNDSDKKEVT 20
   1 PNESYKKOVT 10
   1 ILSEGNSFTA 10
   11 ILSEGNSFTA 20
  FEATURE:
NAME/KEY: MISC_FEATURE
   linear
   09-142-524D-69
  US-09-142-524D-69
  TOPOLOGY:
   US-08-467-023-51
  SEQ ID NO 69
LENGTH: 15
  TYPE: PRT
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Sequence 29, Application US/09014416

Sequence 29, Application US/09014416

Sequence 29, Application US/09014416

Sequence 21, Application US/09014416

Sequence 20, Application US/09014416

APPLICANT: Bukh, Jens

APPLICANT: Bukh, Jens

TITLE OF INVENTION: USES THEREOF

TITLE OF INVENTION: USES THEREOF

TITLE OF INVENTION: USES THEREOF

CURRENT APPLICATION NUMBER: US/09/014,416

CURRENT FILING DATE: 1998-01-27

EARLIER PILING DATE: 1998-01-27

SAFLIER PILING DATE: 1997-07-18

NUMBER OF SEQ ID NOS: 65

SOFTWARE: Patentin Ver. 2.1

LENGTH: 19
   WS-09-014-416-31

WS-09-014-416-31

Sequence 31, Application US/09014416

Patent No. 6153421

GENERAL INFORMATION:

APPLICANT: Bukh, Jens

APPLICANT: Bukh, Jens

APPLICANT: Bukh, Jens

TITLE OF INVENTION: CLONED GENOMES OF INFECTIOUS HEPATITIS C VIRUSES AND

TITLE OF INVENTION: USES THEREOF

TITLE OF INVENTION: USES THEREOF

TITLE OF INVENTION: USES THEREOF

CURRENT APPLICATION NUMBER: US/09/014,416

CURRENT APPLICATION NUMBER: US 60/053,062

EARLIER FILING DATE: 1998-01-27

BARLIER FILING DATE: 1997-07-18

NUMBER OF SEQ ID NOS: 65

SOFTWARE: Patentin Ver: 2.1

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Pred. No. 53;
4; Mismatches 2; Indels
   Length 19;
                         Indels
  Indels
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   Score 34; DB 3;
Pred. No. 53;
  33.3%; Pred. No. 43; tive 7; Mismatches
   Pred. No. 53;
4; Mismatches
  33.7%;
45.5%;
   33.7%;
45.5%;
   |:|:: | | ::|::
1 SKGSAVTTPQNNDEE 15
  3 SEGNSFTAPNDSDKK 17
   ; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-09-014-416-29
  ORGANISM: Hepatitis C virus
Best Local Similarity 33.3
Matches 5; Conservative
   Query Match 33.7
Best Local Similarity 45.5
Matches 5; Conservative
  Conservative
  6 TYTKPNSSDOR 16
  7 SFTAPNDSDKK 17
  7 SFTAPNDSDKK 17
   :: | | | | ::
6 TYTEPNSSDQR 16
  Query Match
Best Local Similarity
Matches 5; Conserv
   RESULT 10
US-09-014-416-29
   US-09-014-416-31
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   DB 3; Length 19; 36;
   Indels
  COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSET for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,473
FILING DATE: 26-Mar-1998
PRIOR APPLICATION DATA:
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Genencor International, Inc.
STREET: 925 Page Mill Road
CITY: Palo Alto
   4; Mismatches
   APPLICATION NUMBER: 07/214,237
FILING DATE: 01-JUL-1992
APPLICATION NUMBER: 07/931,123
FILING DATE: 17-AUG-1992
APPLICATION NUMBER: 08/345,018
FILING DATE: 23-NOV-1994
   TITLE OF INVENTION: ASPARTIC PROTEINMENTOUS FUNGI
   REFERENCE/DOCKET NUMBER: GC45-4
TELECOMMUNICATION INFORMATION:
              CURRENT APPLICATION NUMBER: US/09/014,416
CURRENT FILING DATE: 1998-01-27
EARLIER PELING DATE: 1997-07-18
EARLIER FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 33
LENGTH: 19
  TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-09-048-473-12
   Score 35;
Pred. No.
   NAME: Glaister, Debra J.
REGISTRATION NUMBER: 33,888
   Sequence 12, Application US/09048473
Patent No. 6509171
GENERAL INFORMATION:
   TELEPHONE: 650-846-7620
   Hayenga, Kirk J.
Lawlis, Virgil B.
Ward, Michael
  ATTORNEY/AGENT INFORMATION:
  LENGTH: 16 amino acids
  ZIP: 94304-1013
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
  APPLICANT: Berka, Randy M.
   Query Match
Best Local Similarity 45.5%;
Matches 5; Conservative
  TYPE: PRT

ORGANISM: Hepatitis C virus

US-09-014-416-33
   NUMBER OF SEQUENCES: 17
   TYPE: amino acid
  ::| || ||::
6 TYTGPNSSDQR 16
   7 SFTAPNDSDKK 17
  FILE REFERENCE: 20264276
   COUNTRY: US
   INFORMATION FOR
  US-09-048-473-12
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33.7%; Score 34; DB 4; Length 16;

Query Match

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Gaps
   ö
  Sequence 170, Application US/08475263
Fatent No. 5759994
GENERAL INFORMATION:
APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave., NW
  DB 1; Length 12; 96;
   4; Indels
  APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVEXTION: RECOMBINANT THROMBIN RECEPTOR AND
TITLE OF INVEXTION: RELATED PHARMACEUTICALS
  MEDIUM TYPE: FILIDDY disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/789,184
FILING DATE: 19911107
CLASSIFICATION: 514
ATTONINY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
   COUNTRY: USA
ZIP: 2006-1812
COMPUTER RADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,263
  REFERENCE/DOCKET NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 22000-20502.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: 24.0792
   1; Mismatches
   Score 31;
Pred. No.
Sequence 170, Application US/07789184 Patent No. 5688768 GENERAL INFORMATION:
  E: MORRISON & FOERSTER
755 Page Mill Road
  TELERAX: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 170:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
   30.7%;
58.3%;
  Query Match
Best Local Similarity 58.3
Matches 7; Conservative
   7 SFTAPNDSDKKE 18
  1 SFLARNPNDKYE 12
   NUMBER OF SEQUENCES: 2:
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON
  COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
  CITY: Palo Alto
STATE: California
   ; TOPOLOGY: linear
US-07-789-184-170
  RESULT 15
US-08-475-263-170
  STREET:
   STATE:
   ò
  엄
   APPLICANT: Kume, Akinori
APPLICANT: Kume, Akinori
APPLICANT: Dairiki, Kazuo
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Akiko
APPLICANT: Ivama, Akiko
APPLICANT: Kino, Kohsuke
TITLE OF INVENTION: Peptide-Based Immunotherapeutic Agent for Treating Allergic Disea
FILE REFERENCE: SPO-103
CURRENT APPLICATION NUMBER: US/09/142,524D
PRIOR APPLICATION NUMBER: PCT/JP97/00740
PRIOR APPLICATION NUMBER: PCT/JP97/00740
NUMBER OF SEQ ID NOS: 174
SOFTWARE: Patentin version 3.1
                                   US-09-014-416-35
Sequence 35, Application US/09014416
Patent No. 6153421
GENERAL INFORMATION:
APPLICANT: Vanagi, Masayuki
APPLICANT: Bukh, Jens
APPLICANT: Bukh, Jens
APPLICANT: Bukh, Jens
APPLICANT: Bukh, Jens
TITLE OF INVENTION: CLONED GENOMES OF INFECTIOUS HEPATITIS C VIRUSES AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 20264276
CURRENT APPLICATION NUMBER: US/09/014,416
CURRENT FILING DATE: 1998-01-27
EARLIER FILING DATE: 1998-01-27
EARLIER FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn Ver. 2.1
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   Score 34; DB 3; Length 19;
Pred. No. 53;
4; Mismatches 2; Indels
   Score 33; DB 4; Length 15;
Pred. No. 59;
0; Mismatches 3; Indels
  LOCATION: (1). (15)
COTHER INFORMATION: Cryj2 peptide, Figure 2, Row 34
US-09-142-524D-117
   Sequence 117, Application US/09142524D Patent No. 6719976
  ORGANISM: Cryptomeria japonica
   32.7%;
70.0%;
   33.7%;
  ; ORGANISM: Hepatitis C virus US-09-014-416-35
   Query Match
Best Local Similarity 70.0
Matches 7; Conservative
   Query Match 33.7
Best Local Similarity 45.5
Matches 5; Conservative
  7 SFTAPNDSDKK 17
  6 AYTEPNSSDOR 16
  5 GNSFTAPNDS 14
  2 GISITAPRDS 11
  APPLICANT: Sone, Toshio
   NAME/KEY: MISC_FEATURE
   GENERAL INFORMATION:
   -09-142-524D-117
   RESULT 14
US-07-789-184-170
   SEQ ID NO 35
   SEQ ID NO 117
LENGTH: 15
  TYPE: PRT
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